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(revised) (first entry)

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ABG94320 ID ABG9 XX ABG9 XX ABG9 XX ABG9 XX PQB2 XX PQB2 XX PQB2 XX Huma XM CYtc XX Unid XX Un Renner WA, Piossek C; Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory; cytostatic; antiviral; antidiabetic; hypoglycaemic; antigen array; vaccine; infectious disease. WPI; 19-JAN-2001; 2001US-0262379P. 04-MAY-2001; 2001US-0288549P. 05-OCT-2001; 2001US-0326998P. 07-NOV-2001; 2001US-0331045P. Molecular 21-JAN-2002; 2002WO-IB000166 25-JUL-2002 WO200256905-A2 unidentified bacteriophage. (CYTO-) 2002-627351/67. CYTOS antigen Bachmann M, BIOTECHNOLOGY AG. array used Tissot ä the þ production Maurer ٦, of f Lechner vaccines Ή, for infectious Sebbel P

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This invention relates to a novel ordered and repetitive antigen array used in the production of vaccines for infectious diseases. The invention also discloses a composition comprising a non-natural molecular scaffold comprising a core particle selected from a core particle of a non-natural origin and a core particle of natural origin and an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond. Also disclosed is an antigen or antigenic determinant with at least one second attachment

Claim diseases.

18;

Page

426;

441pp;

English.

ABG94316 ABG80628

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RESULT 2
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05-OCT-2001;
07-NOV-2001;
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CC through the association to form an ordered and repetitive antigen array.

CR Also included is a process for producing a non-naturally occurring ordered and repetitive antigen array. The composition is used in immunisation and as a vaccine for diseases such as influenza, graft crespiratory distress syndrome (ARDS), Crohn's disease, allergic asthma, respiratory distress syndrome (ARDS), Crohn's disease, allergic asthma, creations, anaphylaxis, adult crespiratory distress syndrome (ARDS), Crohn's disease, allergic asthma, creations, and creations, anaphylaxis, adult creations, industriations, influenza, graft creations, influenza, disease, graft creations in figure disease, multiple sclerosis, myasthenia creations, influenza, disease, multiple sclerosis, Alzheimer's disease, creations, and infectious diseases. The antigens are modified to posses a cleavage site (enterokinase or factor Xa) and a Cysteine-containing Noton C-terminal linker peptide which serves as the attachment point to a present sequence is bacterial protein (the scaffold protein). The present sequence is bacterial protein or peptide which is coupled to the modified antigen to form the molecular antigen array
                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a composition comprising: (a) a non-natural molecular scaffold comprising: (i) a core particle selected from: (l) a core particle of a non-natural origin; and (2) a core particle of a non-natural origin; and (ii) an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one second attachment site, where the antigenic determinant with at least one second attachment site, where the antigen or antigenic determinant is amyloid beta peptide (Abeta 1-42) or its fragment, and where the second attachment site is selected from: (i) an attachment site not naturally occurring with the antigen or antigenic determinant; and (ii) an attachment site naturally occurring with the antigen or antigenic determinant; and of association through at least one non-peptide bond to the first attachment site; and where the association of antigenic determinant and the scaffold interact
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 18; Page 146;
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LIDAIDQLNPAY 132
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                                                                                                                                                 Score 664; DB 5;
Pred. No. 2.3e-71;
); Mismatches 0;
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RESULT 3 ABR56455 ID ABR5

ABR56455

standard; protein; 132

cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory;

Antigen presenting cell; APC; immune response; virus like particle;

VLP;

S 밁 ð. B

밁

Bacteriophage Q-beta coat protein mutant SEQ ID NO:26

(first entry)

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The present invention describes a composition (C) for enhancing an immune CC cresponse against an antigen or a virus-like particle (VLP) bound to at least one antimal. (C) CC comprises a virus-like particle (VLP) bound to at least one antimal. Also CC vLP capable of being recognised by the immune system of the animal. Also CC described: (1) enhancing an immune response against an antigen or a VLP in an animal comprising introducing (C) into the animal; (2) vaccines comprising (C) together with a pharmaceutical diluent, carrier or cexcipient; (3) immunising or treating an animal comprising administering the vaccine; and (4) enhancing anti-viral protection in an animal comprising introducing (C) into the animal. (C) compressive, antiaddictive, antiparastic, fungicide, antiallergic, immunosuppressive, antiaddictive, antianflammatory, cantiallergic, immunosuppressive, antiaddictive, antificammatory, cantitheumatic and antiarthritic activities. (C) or the vaccines can be used for enhancing antimune response against an antigen or a VLP in an animal, enhancing anti-viral protection in an animal, or immunising or treating tumours and infectious diseases such as viral, bacterial, continuation or fungal infectious diseases such as viral, bacterial, continuation or treating altergies, drug addiction, graft-versus-host of parasitic or fungal infectious disease, drug addiction, graft-versus-host content to the state of the present earnerse. Accessas to Accessas to Accessas and ABR5401 to ABR56509
Query Match
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Matches 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A composition, useful for enhancing an immune response against an antig or a virus-like particle, enhancing anti-viral protection in an animal, or immunizing or treating tumors or infectious diseases, e.g. viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 182; 243pp; English
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Score 664; DB 6;
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Best Local Similarity

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Score 664; DB 6; Pred. No. 2.3e-71;

Length 132;

Query Match

Sequence 132

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The invention describes an angiotensin peptide moiety carrier conjugate comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide moiety with at least one second comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kidney failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta coat protein used in the preparation of the vaccine conjugates of the invention
                                                                                                                                                                                                                                                                                                                       New angiotensin peptide moiety carrier conjugate comprising a carrier and an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-OCT-2001; 2001US-0326998P
07-NCY-2001; 2001US-0331045P
10-JAN-2002; 2002US-00059902-
21-JAN-2002; 2002WO-IB000166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteriophage Obeta; coat protein; hypotensive; cerebroprotective; cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction;
                                                                                                                                                                                                                                                                   Claim 20; Page 94; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-430264/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-OCT-2002; 2002WO-EP011219
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        congestive heart failure; kidney failure; retinal haemorrhage; mutant;
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2002US-00050902.
2002WO-IB000166.
2002US-0396637P.
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The present invention describes a composition (C) for enhancing an immune response in an animal. (C) comprises a virus-like particle (VLP), and an immunostimulatory substance. The immunostimulatory substance is bound to the VLP. Also described: (1) enhancing an immune response in an animal by introducing (C) into the animal; (2) producing (C) for enhancing an immune response in an animal; (3) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; and (4) immunising or
                                                                                                                                                                                                                                                                                                                                                                                                                 New compositions comprising immunostimulatory substances packaged virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, turns in the compositions of the composition 
                                                                                                                                                                                                                                                                                                          Disclosure; Page 261-262; 322pp; English.
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(TISS/)
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22-APR-2002; 2002US-0374145P.
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hepatitis B virus; lymphocytic choriomeningitis virus; vaccine;
immunostimulant; cytostatic; antiallergic; virucide; antibacterial;
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MEIJERINK E.

LIPOWSKY G.

PUMPENS P.

CIELENS I.

RENHOFA R.
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Cielens I,
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, Renhofa R,
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, Bachmann MF,
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 132
                                                                                                                                                                                                                                                                                                         18-JAN-2002;
21-JAN-2002;
08-JUL-2002;
                                                                    WPI; 2003-598483/56.
                                                                                                                                    Bachmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vaccine composition, virus-like particle; core particle; first attachment site; antigen; antigenic determinant; prion PrP, PrP peptide; vaccine, neuroprotective, antinflammatory; prion disease; Bovine Spongiform Encephalopathy; BSE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JAN-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003059386-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage Qbeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Creutzfeldt-Jakob Disease; coat protein; mutant; mutein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vaccine composition; virus-like particle;
                                                                                                                                                                                                            (CYTO-)
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                                                                                                                                                                                                         CYTOS BIOTECHNOLOGY AG
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                                                                                                                                                                                                                                                                      ; 2002US-00050902.
; 2002WO-IB000166.
; 2002US-0393725P.
; 2002US-0396590P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      2003WO-EP000460.
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                                                                                                                                    Maurer P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Qbeta coat protein mutant Qbeta-251
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Pred. No. 2.3e-71;
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                                                                                                                                    Renner
                                                                                                                                    WA;
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RRESULT 7
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like or a core particle with at least one first attachment site and at
least one antigen or antigenic determinant that is a prion protein (PrP)
or its dimer, or a PrP peptide, the antigen or antigenic determinant
being bound to the virus-like or core particle. The vaccine of the
invention may have neuroprotective or antiinflammatory activity. The
composition is useful as a medicament or in manufacturing a medicament
for the treatment or prevention of prion diseases. The prion diseases may
include Bovine Spongiform Encephalopathy (BSE) or Creutzfeldt-Jakob
Disease. The present sequence is the amino acid sequence of a mutant coat
protein from bacteriophage Obeta which may be used during the creation of
the vaccine composition of the invention.
                                                                                                                                                                                                                                                                                                    07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050902.
19-JAN-2002; 2002WO-IB000166.
19-JUL-2002; 2002US-0396635P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        phage) and at least one particle.
New compositions comprising a core particle and at least one antigen antigenic determinant, useful for as a vaccine for therapy or prophyl of bone diseases, particularly mammalian encephalopathies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-NOV-2002; 2002WO-EP012449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             osteopathic; vaccine; core particle; antigenic determinant; RANKL; bone disease; encephalopathy; immune system stimulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADJ82059 standard; protein; 132 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Creutzfeldt-Jakob Disease) comprises a virus-like particle (e.g. RNA-phage) and at least one prion protein or peptide bound to the virus-like
                                                                                                                          WPI; 2003-441430/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-MAY-2003.
                                                                                                                                                                                                                                              (CYTO-)
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                                                                                                                                                                                                                                              CYTOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                      Maurer
                                                                                                                                                                                                                                           BIOTECHNOLOGY AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antigen array to treat bone disease
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Pred. No. 2.3e-71;
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RESULT 8
ADK17147
ID ADK11
XX ADK1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 132; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The
                                                                                                                                                                                                                                                                                            07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050906.
21-JAN-2002; 2002WO-IB000166.
19-JUL-2002; 2002US-0396636P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiallergic; antiasthmatic; cytostatic; vaccine; virus-like interleukin; IL-5; IL-13; eotaxin; repetitive antigen array;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 132 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            07-NOV-2002; 2002WO-EP012455.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Virus-like particle repetitive antigen array peptide #26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK17147 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003040164-A2
                                                                                                                                                                                                                                     (CYTO-) CYTOS
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                                                                                                                    2003-441518/41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     132; Conservative
                                                                                                                                                                              Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eosinophilic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                        Jennings
                                                                                                                                                                                                                                     BIOTECHNOLOGY AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide; 132
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                                                                                                                                                                           ຸດ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease; asthma;
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                                                                                                                                                                           Sonderegger I;
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Pred. No. 2.3e-71
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Composition comprising an ordered and repetitive antigen or antigenic determinant array, useful as a medicament, or for manufacturing a medicament for treating allergic eosinophilic diseases, e.g. asthma,

Hodgkin's lymphoma

SEQ ID NO

26; 245pp;

English.

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RESULT 9
ADJ36316
ID ADJ3
AC ADJ3
AC ADJ3
AC Bact
XX ADJ3
AC Bact
XX ADJ3
XX ADJ3
AC Bact
XX ADJ3
XX ADJ3
AC Bact
XX ADJ3
AC Bact
XX ADJ3
AC CATC
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New compositions comprising a virus-like particle (VLP), an immunostimulatory substance bound to the VLP, and an antigen mixed with the VLP, useful for enhancing immune response or for treating e.g. tumo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a composition comprising a virus-like particle and at least one antigen, which is a protein or peptide of interleukin (IL)-5, IL-13 or eotaxin and is bound to the virus-like particle, or a core particle with at least one first attachment site and at least one antigen with at least one second attachment site, where the antigen is a protein or peptide of IL-5, IL-13 or eotaxin. The second attachment site is an attachment site naturally or not naturally occurring with the antigen or antigenic determinant. The second attachment site is capable of association to the first attachment site, and where the antigen or ansignic determinant and the core particle interact through the compositions are useful as medicaments, or for manufacturing a medicament or a vaccine for treating allergic eosinophilic diseases, e.g. asthma, or constraint the communication of the first attachment site states and where the communication to form an ordered and repetitive antigen array. The commositions are useful as medicaments, or for manufacturing a medicament for a vaccine for treating allergic eosinophilic diseases, e.g. asthma, or canararse the common and related diseases. This sequence is used to
                                                                                                                                                                                                                          Bachman MF,
                                                                                                                                                                                                                                                                                                                                                            20-JUN-2002; 2002US-0389898P.
                                                                                                                                                                                                                                                                                                                                                                                                                                20-JUN-2003; 2003WO-EP006541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004000351-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immune response; virus-like particle; immunostimulatory; all chronic disease; chronic viral disease; bacteriophage Qbeta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiallergic; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage
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                                                                                                                                                                                                                                                                                       CYTOS BIOTECHNOLOGY AG
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                                                                                                                                                                                                                       Renner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         compound of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Qbeta coat protein virus-like particle mutant K2RK13R
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Pred. No. 2.3e-71;
Mismatches 0;
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RESULT 10
ADJ677162
ID ADJ677
XX ADJ677
XX DJ677
XX DG-MA)
XX DG-MA)
XX Anore
XX Bacte
OS Synth
XX Bacte
OS Synth
XX ID-JU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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antigenic a
                                                                                                                  Disclosure; SEQ ID NO
                                                                                                                                                                            peptide
                                                                                                                                                                                                                                                                                            WPI; 2004-132866/13.
                                                                                                                                                                                                                                                                                                                                                  Bachmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               an animal comprising a virus-like particle, an immunostimulatory substance bound to the virus-like particle, and an antigen mixed with the virus-like particle, and an antigen mixed with the virus-like particle. The composition or the vaccine is useful in the manufacture of a pharmaceutical for the treatment of a disorder or disease such as allergies, tumours, chronic diseases and chronic viral diseases. The composition is also useful for enhancing an immune response in an animal. This is the amino acid sequence of a bacteriophage Qbeta coat protein mutant, a virus like particle (VLP) that can be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-JUL-2003; 2003WO-EP007849.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUL-2002; 2002US-0396638P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacteriophage Obeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Phage Qbeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADJ67162 standard;
                                                                                                                                                                                                                                                                                                                                                                                                            (CYTO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2004009124-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADJ67162;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 132 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   adjuvant of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 17; 252pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or chronic viral diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a composition for enhancing an immune
                                                                                                                                                             composition comprising a core particle having a first attachment site an antigen or antigenic determinant which is a ghrelin or ghrelin tide having a second attachment site, useful for treating obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           121 LIDAIDQLNPAY 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                                                                                                                                                                                                                                                                                                                            CYTOS BIOTECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                  ΜĘ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AKLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       array.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         core particle; antigenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coat protein mutant Qbeta 251 for antigen display array.
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                                                                                                                                                                                                                                                                                                                                               Fulurija
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                                                                                                                  20;
                                                                                                         175pp;
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Pred. No. 2.3e-71;
Mismatches 0;
                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         determinant; ghrelin; P-pilin;
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The invention relates to a new composition comprising: (i) a core particle with at least one first attachment site; and (ii) at least one antigen or antigenic determinant with at least one second attachment

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     site, where the antigen or antigenic determinant is ghrelin or a ghrelin peptide, and where the second attachment site being consisting of an attachment site not naturally occurring with the antigen or antigenic determinant and an attachment site naturally occurring with the antigen or antigenic determinant, where the second attachment site is capable of association to the first attachment site, and where the ghrelin or a ghrelin peptide and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful for treating obesity. The repetitive array may form part of a phage or bacterial display array. This peptide corresponds to a Bacteriophage obeta coat protein mutant which can used as part of the repetitive or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 132 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuroprotective; nootropic; vaccine; amyloid beta 1-6 peptide;
core particle; antigen array; Alzheimer's disease; RNA bacteriophage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteriophage Qbeta coat protein mutant K16R A88K
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15-MAY-2003; 2003US-0470432P.
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                                                                     WPI; 2004-203731/19.
                                                                                                                                                                   Bachmann MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                      18-JUL-2003; 2003WO-EP007864
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                                                                                                                                                                                                                                      (CYTO-) CYTOS BIOTECHNOLOGY AG. (NOVS) NOVARTIS PHARMA AG.
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                                                                                                                                                                       Tissot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "Wild type Arg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note= "Wild type Lys substituted by Arg"
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                                                                                                                                                                       Ortmann
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Composition comprising a core particle with at least

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AC ABG94
XX ABG94
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Best Local Similarity
      Renner WA,
Piossek C;
                                                                                                                                                       19-JAN-2001; 2001US-0262379P.
04-MAY-2001; 2001US-02885499P.
05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-0331045P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory; cytostatic; antiviral; antidiabetic; hypoglycaemic; antigen array; vaccine; infectious disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 132 AA;
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                                                                                                                                                                                                                                                                                                            21-JAN-2002; 2002WO-IB000166
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10-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABG94319 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200256905-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unidentified bacteriophage.
                                                                                              (CYTO-) CYTOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 LIDAIDQLNPAY 132
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                                  Bachmann M,
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(first en
                                                                                              BIOTECHNOLOGY AG
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                                            Tissot
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Pred. No. 2.3e-71;
); Mismatches 0;
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                                            Maurer
                                         ָס'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     comprising a virus-like core and an antigenic amyloid beta
                                         Lechner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 132;
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                                         শ
                                            Sebbel P;
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ABG80631
ID ABG1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cc site, where the antigen or antigenic determinant is amyloid beta peptide
(Abeta1-42) or its fragment and where the second attachment site is
cc selected from an attachment site not naturally occurring with the antigen
cc or antigenic determinant and an attachment site naturally occurring with
cc the antigen or antigenic determinant, where the second attachment site is
cc capable of association through at least one non-peptide bond to the first
cc tatachment site and where the antigen or antigenic determinant and the
cc scaffold interact through the association to form an ordered and
cc repetitive antigen array. The invention also comprises a coat protein
cc capable of forming a capsid which comprises mutant Obeta coat proteins
cc having an amino acid sequence selected from five amino acid sequences
cf cally defined in the specification. The compounds of the invention may
capable of forming a rapsid which compounds of the invention may
capable of forming a capsid which compounds of the invention may
capable of forming a capsid which compounds of the invention may
capable of forming a capsid which compounds of the invention may
capable of forming a capsid which compounds of the invention may
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cc antiviral, antidabetic, or hypoglycaemic activities and may be used in
cc antiviral, antidabetic, or hypoglycaemic activities and may be
cc antiviral, antidabetic, or hypoglycaemic activities
cc 
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Best Local :
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        molecular scaffold; amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgE-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus erythematosus; osteoporosis; inflammatory immune disease; myasthenia gravis; multiple sclerosis; inmunoproliferative disease; lymphadenopathy; lmmunoblastive lymphadenopathy; angloimmunoproliferative lymphadenopathy; immunoblastive lymphadenopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a novel ordered and repetitive antigen array used in the production of vaccines for infectious diseases. The invention also discloses a composition comprising a non-natural molecular scaffold comprising a core particle selected from a core particle of a non-natural origin and a core particle of natural origin and a core particle of natural origin and a core particle of the comprising at least or first attachment of the control origin and a core particle of the comprising at least or first attachment of the control origin and a core particle of the comprising at least or first attachment of the control origin and a core particle of the comprising at least or first attachment of the control origin and a core particle or the control origin and a core particle or the control origin and a core particle of the control origin and a core particle or the control origin and a core particle origin and a core particle or the control origin and a core particle or
    rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage Q-beta Al coat protein mutant Qbeta-251.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABG80631 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 18; Page 146; 441pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Molecular antigen array used in the production of vaccines for infectious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-627351/67
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| 121 LIDAIDQLNPAY 132
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                                                                                                                                                                                                                                                                                                                                                                    antigen array; vaccine; antigen; antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ç
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Pred. No. 5.3e-71;
1; Mismatches 0
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The invention relates to a composition comprising: (a) a non-natural CC molecular scaffold comprising: (i) a core particle selected from: (1) a CC core particle of a non-natural origin; and (2) a core particle of natural CC origin; and (ii) an organiser comprising at least one first attachment (CC origin; and (ii) an organiser is connected to the core particle by at least one covalent bond; (b) an antigen or antigenic determinant with at least consecond attachment site, where the antigen or antigenic determinant is camploid beta peptide (Abeta 1-42) or its fragment, and where the second attachment site is selected from: (i) an attachment site not naturally occurring with the antigen or antigenic determinant; and (ii) an CC attachment site naturally occurring with the antigen or antigenic core through at least one non-peptide bond to the first attachment site; and through the association to form an ordered and repetitive antigen array. The composition is used in continued is a process for producing a non-naturally occurring cordered and repetitive antigen array. The composition is used in continued as a vaccine for diseases such as influenza, graft corners to be supplied to the seast one analystaxis, adult crespiratory distress syndrome (ARDS), Crohn's disease, allergic asthma, caute lymphoblastic leukaemia, non-Hodgkin's lymphoma, Grave's disease, systemic lupus erythematosus, inflammatory immune diseases, myasthenia corrections, anaphylaxis, adult crespiratory differative disease lymphadenopathy.
                               angioimmunoproliferative lymphadenopathy, immunoblastive lymphadenopathy, rheumatoid arthritis, diabetes, multiple sclerosis, Alzhaimer's disease, osteoporosis and infectious diseases. The antigens are modified to posses a cleavage site (enterokinase or factor Xa) and a Cysteine-containing Nor C-terminal linker peptide which serves as the attachment point to a virus like particle or bacterial protein (the scaffold protein). The present sequence is bacterial protein or peptide which is coupled to the modified antigen to form the molecular antigen array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Maurer
Renner
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04-MAY-2001; 2001US-0288549P.
05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-0331045P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 18; Page 403; 418pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Molecular antigen array used in the
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ORTMANN
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Tissot A,
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Sebbel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       production of vaccines for infectious
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Matches 131; Best Local Similarity

Conservative

1;

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Gaps

0

99.5%;

Score 661; DB 5; Pred. No. 5.3e-71; Mismatches

Length 132;

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RESULT 14
ABR56456
ID ABR56
XX ABR56
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                              The present invention describes a composition (C) for enhancing an immune cresponse against an antigen or a virus-like particle in an antigen, or a virus-like particle (VLP) bound to at least one antigen, or a VLP capable of being recognised by the immune system of the animal. Also described: (1) enhancing an immune response against an artigen or a VLP in an animal comprising introducing (C) into the animal; (2) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; (3) immunising or treating an animal comprising administering the vaccine to the animal, or priming or boosting a T cell response in the animal by administering the vaccine; and (4) enhancing anti-viral protection in an animal comprising introducing (C) into the animal. (C) has cytostatic, virucide, antibacterial, antiparasitic, fungicide, antitallergic, immunosuppressive, antiaddictive, antiinflammatory, antitinflammatory, antitinflammat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory; antithyroid; antidiabetic; neuroprotective; nootropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis; Alzheimer's disease; osteoporosis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A composition, useful for enhancing or a virus-like particle, enhancing or immunizing or treating tumors or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 183; 243pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003024480-A2
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    an antigen
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Matches 131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-IB000166.
19-JUL-2002; 2002US-0396637P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction; congestive heart failure; kidney failure; retinal haemorrhage; mutant
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New angiotensin peptide moiety carrier conjugate comprising an angiotensin peptide moiety, useful for treating or prever disorder associated with renin-activated angiotensin, e.g. h

preventing e.g. hyper

2003-430264/40.

Claim 20;

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English.

describes an angiotensin peptide moiety carrier conjugate

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Search completed: January 4, 2005, 09:11:02 Job time: 40.6824 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kichey failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta coat protein used in the preparation of the vaccine conjugates of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 132 AA;
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/cgm2_6/ptodata/1/iaa/6A_COMB.pep:*
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      GenCore version (c) 1993 - 2005
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US-09-200-673-15
US-09-200-673-15
US-10-013-846-4
US-09-200-673-15
US-10-013-846-4
US-09-200-673-15
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US-09-200-673-15
US-08-725-4598-45
US-08-725-4598-44
US-08-725-4598-44
US-08-725-4598-44
US-08-725-4598-44
US-08-725-4598-42
US-09-216-9148-1
US-09-218-990-148-1
US-09-218-992-19
US-09-248-796A-16565
US-09-248-796A-16565
US-09-38-239-5
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Compugen Ltd.
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Sequence 15, Appli
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Sequence 44, Appli
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Sequence 47, Appli
Sequence 1650, Appli
Sequence 17, Appli
Sequence 216, Appli
Sequence 5, Appli
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RESULT 2 US-09-948-722-2

Sequence 2, Application US/09948722
Patent No. 6776993
GENERAL INFORMATION:
APPLICANT: Kaufmann, Stefan H. B.
APPLICANT: Hess, Jurgen
TITLE OF INVENTION: Tuberculosis Vaccine

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TLSIDLPGMTNQDNKIVVK--NATKSNVNNAVNTLVERWNEKYAQAYPNVS---AKIDYD

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49 TV	13 KD	atch cal	FEATURE: OTHER INFORMATION: 09-485-717-2	PRT ISM:	N	APPL	R APPLICATION NUMBER: R FILING DATE: 1997-08	ENT APPLI	TITLE OF INVENT	CANT:	-71; e 2		66	, 6		٠,٠	67 7	67.5	١	6 6 60 60	68	თ თ თ	6 8	68.5	
SVSQPSRI	KDGRQTLVL- : : : KDGNEYIVVE	Similarity O; Conservat		e Artificial	PatentIn Ver		TION NUT	APPLICATION NUMBER: FILING DATE: 2000-	ION:	RMATION: Kaufmann,	7-2 , Applicatio		9.9				•		10.2				•	٠	
RKOYKVQVI	SKKKKSIN	12.7 % 22.7 % /ative	Description	l Sequence	er. 2.1	NUMBER: PC: 1998-08-1	ПМВЕR: EP 1997-08-22	Ţ.:-	Tubercul	Stefan	on US/09485717		1281 4	~					936 4					1065 4	
TVSVSOPSRNRKNYKVQVKIQNPIACTANGSCDPSVTR	KDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRV	; Score 84.5; DB 4; ; Pred. No. 0.16; 23; Mismatches 44;	tion of Artificial Sequence:	nce		PCT/EP98-05109 -12	971146	9/	losis Vaccine		9485717	ALIGNMENTS	US-08-843-530B-6 US-09-636-728-5	US-09-134-001C-	US-09-1/3-581-7	US-08-397-411-13	SUS	US S	US-08	US-09-421-868-2 US-08-956-171E-524	US-08	US-08-856-253-7	US-09-270-767-	US-09-560-876A-6	
QKYADVTFSFTQYSTD		Length 626; Indels 35; Gaps	nce: recombinant										Sequence 5, Appli	Sequence 2885,	Sequence 7, App		Sequence 41, Ap	Sequence 76, Ap	Sequence 5249, Ap	Sequence 5249,	Sequence 2, Appli	Sequence 7, App	4 0	e 6,	
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                                                                                                                                                                                                                                                                                           ; ORGANISM: Acinetobacter baumannii
US-09-328-352-4249
                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
SEQ ID NO 4249
                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 4249, App
Patent No. 6562958
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Conser
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CURRENT FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
                                                                                                                                                                                                                                                                                                                               LENGTH: 40
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: EP 97/114,614.7 PRIOR FILING DATE: 1997-08-23 NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/485,717
PRIOR PILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: PCT/EP98/05109
PRIOR FILING DATE: 1998-08-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/948,722
CURRENT FILING DATE: 2002-04-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 100564-00079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
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LENGTH: 626
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214 ASKQILDDLPQL 225
                                117 ASPLLIDAIDOL 128
                                                                  164 RTGSN----AIAYLRETGFTNNAALVPENTAKPYSEITF-
                                                                                                                                     109 SLINFTKOVRGGTRLNVNVPRNALISFAVNPVDGTTPI----IAKPNQRLTIRDLLAPG 163
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                                                                                               57 RNRKNYKVQVKIQNPTACTANGSCDPSVTRQKYADVTPSFTQYSTDEERAFVRTELAALL 116
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                                                                                                                                                                                                                   l Similarity
35; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TLSIDLPGMTNQDNKIVVK--NATKSNVNNAVNTLVERWNEKYAQAYPNVS---AKIDYD 252
                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/09328352
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description of Artificial Sequence: recombinant nucleic acid molecule comprising a domain of Mycobacterium and a phagolysomal
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Pred. No. 0.16;
23; Mismatches 44;
                                                                                                                                                                                                                                  Score 79.5; DB Pred. No. 0.35;
                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                 DB 4;
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LENGTH: 916
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: MAIC J. RUBENFIELD et al.
APPLICANT: MAIC J. RUBENFIELD et al.
TITLE OF INVENTION: AUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                       TELEFAX: (317) 276-3861
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 383 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 6
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Pseudomonas aeruginosa
                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (317) 276-0756
                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: GAYLO, PAUL J.
REGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: P-
                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Baez, Melvyn
APPLICANT: Cates, Carolyn A.
TITLE OF INVENTION: RHESUS NEUROPEPTIDE Y1 RECEPTOR
    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                          TYPE:
                       TOPOLOGY:
                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    56 SRNRKNYKVOVKIONPTAC---TANGSCDPSVTROKYADVTFSFTOYSTDEERAFVRT 110
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                                      amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     United States of America
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protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11.2%; Score 74.5; D
23.7%; Pred. No. 5.3;
                                                                                                                                                                                                                                                                                                      US/09/045,186
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                                                                                                                                                                                    P-11376
                                                                                                                                                                                                                                                                                                                                            Version
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RESULT 7
US-08-555-268A-15
; Sequence 15, Application US/08555268A
; Patent No. 5958709
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US-08-232-144-4
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                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TREEPHONE: 202-783-6040
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Human Neuropeptide Y-Y1 Receptor NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rothwell, Figg, Ernst & Kurz
STERET: 555 13th St, N.W., Suite 701-East
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: ERNST, Barbara G
REGISTRATION NUMBER: 30,377
REFERENCE/DOCKET NUMBER: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                        TOPOLOGY: 11
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 26-MAY CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
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                                                                                                                                                                         138 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN------
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5. 5571695
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                                                                                                                                                                                                16 RQTLVLNPRGVNPTN-----GVA----SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
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                                                                                                                                         68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
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HERZOG, Herbert
SHINE, John
                                                                                                                                                                                                                                                                                                                                                                                    384 amino acids
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25.9%; Pred. No. 3
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                                                                                                          VTLDAYKDKYVCFDQFPSDSHRLSYTTLLLVL 218
                                                                                                                                                                                                                                     ; Score 71.5; E; Pred. No. 3.2; 11; Mismatches
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; MOLECULE TYPE:
US-08-555-268A-15
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                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                             Sequence 15, Appli
Patent No. 6316203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                   APPLICANT: Gerald, Christophe P.G.
APPLICANT: Weinshank, Richard L.
APPLICANT: Walker, Mary W.
APPLICANT: Walker, Mary W.
APPLICANT: Branchek, Theresa
TITLE OF INVENTION: Methods of Modifying Feeding Behavior, Compounds
TITLE OF INVENTION: in Such Methods, and DNA Encoding A Hypothalamic
TITLE OF INVENTION: Atypical Neuropeptide Y/Peptide YY Receptor (Y5)
FILE REFERENCE: 46166-BZ/JDW
CURRENT APPLICATION NUMBER: US/09/200,673A
CURRENT FILING DATE: 1998-11-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (212) 391-0526
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 4474
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy of COMPUTER: IBM PC com
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            138 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN------
FILING DATE: 1994-12-02
                                  APPLICATION NUMBER: 08/566,096 FILING DATE: 1995-12-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16 RQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
                  APPLICATION NUMBER: 08/349,025
                                                                                                                                                                                                                                                                                                                                                                                                                                                   68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
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Y: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                  -------VTLDAYKDKYVCFDQFPSDSHRLSYTTLLLVL 218
                                                                                                                                                                                                                                                                                                                  Application US/09200673A
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1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     384 amino acids
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VENTION: DNA ENCODING A HUMAN NEUROPEPTIDE

VENTION: Y/PEPTIDE YY/PANCREATIC POLYPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10.8%; Score 71.5; Di 25.9%; Pred. No. 3.2;
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                                                                                                                                                                          Compounds Useful
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RESULT 10
US-09-708-392-9
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US-10-013-846-4
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US-09-200-673-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 384
                                                       GENERAL INFORMATION:
                                                                         Sequence 9, Application US/09708392
Patent No. 6734186
                                                                                                                                                                                                                                                                                                                                                 Query Match 10.8%;
Best Local Similarity 25.9%;
Matches 28; Conservative 11
              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/254,990 PRIOR FILING DATE: 2000-12-12 NUMBER OF OF OTHER PRIOR FILING DATE: 2000-12-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Zheng, Xiaozhang
APPLICANT: Elliott, Richard L
APPLICANT: Hammond, Marlys
TITLE OF INVENTION: Spiro(isobenzofuran-1,4'-piperidin]-3-ones and
TITLE OF INVENTION: 3H-spiroisobenzofuran-1,4'-piperidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE: N00.2001
ENT APPLICATION NUMBER: US/10/013,846
ENT FILING DATE: 2001-12-11
                                                                                                                                                                                            138 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4, Application US/10013846
o. 6566367
                                                                                                                                                                                                                               68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
                                                                                                                                                                                                                                                                                                            16 RQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
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                Pfizer Inc
Pfizer Limited
Wayman, Chris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hutchison, Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Brielmann, Harry L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tran, Jennifer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       De Lombaert,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Blum, Charles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ----VTLDAYKDKYVCFDQFPSDSHRLSYTTLLLVL 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stephane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rajagopal
                                                                                                                                                                                                                                                                                                                                                 11;
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                                                                                                                                                                                                                                                                                                                                                 Score 71.5; DB
Pred. No. 3.2;
11; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                        DB 4; Length 384;
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                                                                                                                                                                                                                                                                                                                                                   34; Indels 35;
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RESULT 11
PCT-US93-05039-3
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; ORGANISM: Homo sapiens
US-09-708-392-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE:
SEQ ID NO 9
LENGTH: 38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                              COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: MS-DOS
SOFTWARE: Microsoft Word 4.0
CURRENT APPLICATION DATA:
APPLICATION UNMBER: PCT/US93/05039
FILING DATE: 19930527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Pharmaco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2000-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 2000-(
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/708,392
CURRENT FILING DATE: 2000-11-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
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CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                          APPLICANT: Claes R. TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: GB (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16 RQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
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                                                                                                                                                                                                                                                                                                                                                                                                             Application PC/TUS9305039
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                                                                                                                                                                                                      Connecticut
                                                                                                                                                                                                                                     E: Yahwak & Associates
25 Skytop Drive
                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                                                                                                          Claes R. Wahlestedt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -----VTLDAYKDKYVCFDQFPSDSHRLSYTTLLLVL 218
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2000-01-07
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                                                                                                                                                                                                                                                                                                                    Human Neuropeptide Y/Peptide Receptor of the Y1-Type and Anitsense Oligonucleotides
                                                                                                                                                                                                                                                                                                          Thereto Which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10.8%; Score 71.5; DB 4; Length 384; 25.9%; Pred. No. 3.2;
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                                                                                                                                                                                                                                                                                                          Inhibit Vasoconstriction
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REFERENCE/DOCKET NUMBER: CR TELECOMMUNICATION INFORMATION:

REGISTRATION NUMBER:

26,824

CRF D-1250

TELEPHONE:

(203) 268-1951

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(203) 268-1951

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RESULT 12
US-08-817-869-3
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Best Local Similarity
Matches 28; Conserv
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APPLICANT: STRADE
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                                                                          TELEFAX: (732) 594-4720 INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                         REFERENCE/DOCKET NUMBER: 19339Y PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (732) 594-1348
                                                                                                                                                                       FILING DATE: 07-NOV-1994
ATTORNEY/AGENT INFORMATION:
NAME: NICHOLSON, WILLIAM H.
REGISTRATION NUMBER: 25,147
                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 MOLECULE TYPE:
                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: AMINO ACID
                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                          FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 07065-0900
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                             STRANDEDNESS:
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                   TOPOLOGY:
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                                                amino acid
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                                                             411 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  126 EAST LINCOLN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   384 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MACNEIL, DOUGLAS J.

TVENTION: MODIFIED NEUROPEPTIDE Y RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                   PatentIn Release #1.0, Version #1.30
                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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peptide
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                             single
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                                                                                                                                                                       25,147
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Pred. No. 3.2;
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Query Match

10.8%;

Score 71.5;

DB 3;

Length 411;

US-08-725-459B-45; Sequence 45, Application US/08725459B; Patent No. 6084068

RESULT 14

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PCT-US95-14377-3
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; MOLECULE TYPE:
PCT-US95-14377-3
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                                                                                                                                     Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 07065-0900
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FORM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #11.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: APPOLLINA, MARY A.
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 07-NOV-1994
                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
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CORRESPONDENCE ADDRESS:
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CITY: RAHWAY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
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                                                                                                                                                                                                                                                                               LENGTH:
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     190
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                                                                   141 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN---
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                                     68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTBLAAL 115
                                                                                                   16 ROTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
                                                                                                                                    28;
                                                                                                                                                                                                                                                                 amino acid
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CASCIERI, MARGARET A.
MACNEIL, DOUGLAS J.
                                                                                                                                                                                                                                                                               411 amino acids
                                                                                                                                                                                                                                                                                                                                 (908) 594-4720
                                                                                                                                    Conservative
                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                  (806)
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                                                                                                                                                   10.8%;
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VTLDAYKDKYVCFDQFPSDSHRLSYTTLLLVL 221
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                                                                                                                                  11; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                    19339Y PCT
                                                                                                                                                 Score 71.5;
Pred. No. 3.
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GENERAL INFORMATION:

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US-08-725-459B-44
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Matches
                                                                                                                                                                                                                                                         Sequence 44, Application US/08725459B Patent No. 6084068 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 214-981-3400
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
COUNTRY: US
COUNTRY: US
ZIP: 75201-6507
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: HANSEN, EUGENIA S.
REGISTRATION NUMBER: 31,966
REFERENCE/DOCKET NUMBER: 111
TELECOMMUNICATION INFORMATION:
TELEPHONE: 214-981-3300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/725,459B FILING DATE: 04-OCT-1996 CLASSIFICATION: 530 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: SIDLEY & AUSTIN
STREET: 717 N. HARWOOD, SUITE 3400
                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: SIDLEY & AUSTIN
                                                                                                                                                                                   APPLICANT: CONAWAY, RONALD C.
APPLICANT: CONAWAY, JOAN W.
TITLE OF INVENTION: ELONGIN A AND C FUNCTIONAL DOMAINS
NUMBER OF SEQUENCES: 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICANT: CONAWAY, JOAN W.
TITLE OF INVENTION: ELONGIN A AND C FUNCTIONAL DOMAINS
NUMBER OF SEQUENCES: 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
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OTHER INFORMATION:
OTHER INFORMATION:
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ZIP: 75201-6507
                                                                                    STATE:
                                                                                                         CITY:
                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: not rele
TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                     171 QLSQARKNVKIEGKAQLRT-ITPRGGGVPSTSRSR 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 116 KLEMLT-SRIGKSNSAQSQGRQTMVIDMAHTR----VRSKSFFNTVKDSQVKMSATPSAL 170
                                                                                                                                                                                                                                                                                                                                                                                                                                          54 QPSRNRKNYKVQVKIQNPTACTANGSCDPSVTRQK 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 KLETVTLGNIGK-----DGRQTLVLNPRGVNPTNGVASLSQAGAV--PALEKRVTVSVS 53
                                                                                  DALLAS
: TX
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: TX
                                                                                                                       717 N. HARWOOD, SUITE 3400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "amino acids 202-434 of C. elegans"
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/725 ***
FILING DATE: 04-00**
CTASSIET***: 04-00**
                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: HANSEN, EUGENIA S.
REGISTRATION NUMBER: 31,966
REFERENCE/DOCKET NUMBER: 111
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 341 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                       NAME/KEY: Peptide LOCATION: 1.341 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: not
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279 QLSQARKNVKIEGKAQLRT-ITPRGGGVPSTSRSR 312
                                                              224 KLEMLT-SRIGKSNSAQSQGRQTMVIDMAHTR----VRSKSFFNTVKDSQVKMSATPSAL 278
                              54 OPSRNRKNYKVOVKIONPTACTANGSCDPSVTROK 88
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                                                                                           2 KLETVTLGNIGK-----DGRQTLVLNPRGVNPTNGVASLSQAGAV--PALEKRVTVSVS 53
                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                      341 amino acids
                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                        peptide
                                                                                                                                            10.7%; Score 71;
33.7%; Pred. No.
                                                                                                                                                                                                       /note= "amino acids 94-434 of elegans"
                                                                                                                           11;
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                                                                                                                              Mismatches
                                                                                                                                                           DB 3;
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Search completed: January 4, 2005, 09:35:45
Job time : 11.3622 secs

Copyright

GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.

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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Title:
Perfect score:
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664
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-6/ptodata/1/pubpaa/USO9A_PUBCOMB.pep:*
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_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0	Query
132 1332 1332 1332 1332 1332 1332	Length
144 144 155 144 144	BB
US-10-243-739-26 US-10-284-065-26 US-10-289-458-26 US-10-050-902-259 US-10-346-190-26 US-10-346-190-26 US-10-289-456-26 US-10-622-064-9 US-10-622-064-9 US-10-622-087-20 US-10-622-087-20 US-10-622-087-20 US-10-243-739-27 US-10-244-065-27	ID
Sequence 26, Appl Sequence 26, Appl Sequence 25, Appl Sequence 259, App Sequence 26, Appl Sequence 26, Appl Sequence 26, Appl Sequence 26, Appl Sequence 20, Appl Sequence 20, Appl Sequence 20, Appl Sequence 20, Appl Sequence 27, Appl	Description

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132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	1 + 1
14	14	14	14	14	14	14	14	14	14	14	14	16	15	15	15	15	14	14	14	14	14	14	16	15	15	15	15	14	14	14	
US-10-050-902-257	US-10-050-902-256	-10-050-902-15	54-	54-	89-454	0-244-065-	0-244-065-	44-065-	0-243-739	43-739-	39.	0-622-0	-622-124-1	-622-	-10-289-456-	-465-	-10-346-190-	US-10-050-898-255	0-050-902-	-10-289-454-2	065-	-10-243-739-2	-10-622-087-	-10-622-124-	-10-622-	89-456-	-465-811-	-10-346-190-2	0-050-898-25	-10-050-90	1
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•	256, App											17, Appl	17, Appl		23, Appl		23, Appl		255, App	•	•	23, Appl	•	•	•		18, Appl		_	258, App	,

ALIGNMENTS

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/ ORGANISM: Bacteriophage Q-beta
US-10-243-739-26
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 26
                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                   Matches 132;
                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILLING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
                                                                                                                                                                                                                                          LENGTH: 132
TYPE: PRT
 121 LIDAIDQLNPAY 132
                                                  61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
                               61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
                                                                                                         1 AKLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60
                                                                                           AKLETYTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60
                                                                                                                                                  100.0%; Score 664; DB 14; ilarity 100.0%; Pred. No. 1.6e-66; Conservative 0; Mismatches 0;
                                                                                                                                                                                 Length 132;
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CURRENT APPLICATION NUMBER: US/10/244,065
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR FILING DATE: 2002-04-22
PRIOR APPLICATION NUMBER: 60/318,994
PRIOR FILING DATE: 2001-09-14
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOPTWARE: PATENTIN VERSION 3.1
SEQ ID NO 26
LENGTH: 132
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equence 26, Application US/10289454
publication No. US20030157479A1
GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR FILING DATE: 2002-01-18
PRIOR FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
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Matches
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APPLICANT: Storni, Ta
APPLICANT: Maurer, Pr
APPLICANT: Tissot, Al
APPLICANT: Schwarz, I
APPLICANT: Medjerink
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-10-244-065-26
                                                                                                                                                                                                                              APPLICANT: Bachmann, Martin
APPLICANT: Jennings, Gary
APPLICANT: Jennings, Gary
APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
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APPLICANT: Meijerink, Edwin
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Pumpens, Paul
APPLICANT: Cielens, Indulis
APPLICANT: Renhofa, Regina
APPLICANT: Renhofa, Regina
TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Particl
TITLE OF INVENTION: Method of Preparation and Use
TILE REFERENCE: 1700, 0220001
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mes 132; Conserv
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Maurer, Patrick
Tissot, Alain
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vo. US20030099668A1
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100.0%; Pred. No. 1.6e-66;
tive 0; Mismatches 0;
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APPLICANT: Piossek, Christine
TITLE OF INVENTION: Molecular Antigen Array
FILE REPERENCE: 1700.0190004
CURRENT APPLICATION NUMBER: US/10/050,902
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-05-04
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TYPE: PRT
; ORGANISM: Bacteriophage Q-beta-251
US-10-289-454-26
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US-10-050-902-259
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SEQ ID NO 259
LENGTH: 132
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 26
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/331,045 PRIOR FILING DATE: 2001-11-07
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les 132; Conserv
121
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Lechner, Franziska
Sebbel, Peter
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LIDAIDQLNPAY 132
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Tissot, Alain
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o. US20030175290A1
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100.0%; Pred. No. 1.6e-66;
htive 0; Mismatches 0;
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100.0%; Pred. No. 1.6e-66;
tive 0; Mismatches 0;
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GENERAL INFORMATION:
APPLICANT: Bachmann, Martin
APPLICANT: Maurer, Patrick
APPLICANT: Pellicioli, Erica
APPLICANT: Pellicioli, Erica
APPLICANT: Renner, Wolfgang A.
FILE REFERENCE: 1700.0290003
FILE REFERENCE: 1700.0290003
CURRENT APPLICATION NUMBER: US/10/346,190
CURRENT FILING DATE: 2003-01-17
FRIOR APPLICATION NUMBER: 60/396,590
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR FILING DATE: 2002-07-18
FRIOR APPLICATION NUMBER: 60/398,898
PRIOR APPLICATION NUMBER: 60/389,898
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PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR PILING DATE: 2001-0-05
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR PILING DATE: 2001-11-07
NUMBER: OF SEQ ID NOS: 350
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 259
LENGTH: 132
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; ORGANISM: Qb 251
US-10-050-898-259
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US-10-346-190-26
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US-10-050-898-259
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Best Local Similarity
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Sebbel, Peter
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Ortmann, Rainer
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; LENGTH: 132
; TYPE: PRT
; ORGANISM: Qb
US-10-346-190-26
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, ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Bacteriophage Q-beta 251 mutant
US-10-465-811-17
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APPLICANT: RENNER, WOLFGANG A
TITLE OF INVENTION: PACKAGED VIRUS-LIKE PARTICLES FOR USE AS ADJUVANTS:
TITLE OF INVENTION: METHOD OF PREPARATION AND USE
FILE REFERENCE: 1700.0290004
CURRENT APPLICATION NUMBER: US/10/465,811
CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: US 60/389,898
PRIOR PRILING DATE: 2002-06-20
NUMBER OF SEQ ID NOS: 131
SOPTWARE: Patentin version 3.2
SEQ ID NO 17
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PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR FILING DATE: 2002-01-18
NUMBER OF SEQ ID NOS: 164
SOFTWARE: PatentIn version 3.1
SEQ ID NO 26
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Publication No.
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Best Local Similarity 100.0%;
Matches 132; Conservative 0
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Best Local Similarity
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                       121 LIDAIDQLNPAY 132
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LIDAIDQLNPAY 132
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                                                                                                                                                                                                                                           100.0%; Score 664; DB 15; 100.0%; Pred. No. 1.6e-66;
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Pred. No. 1.6e-66;
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RESULT 8 US-10-289-456-26

; Sequence 26, Application US/10289456

GENERAL INFORMATION:

APPLICANT: Bachmann, Martin APPLICANT: Maurer, Patrick

Publication No. US20040033211A1

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                                                                                                                                                       ; OTHER INFORMATION: Bacteriophage Q-beta 251 mutant
US-10-622-064-9
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CURRENT APPLICATION NUMBER: US/10/289,456

CURRENT APPLICATION NUMBER: US/10/289,456

CURRENT FILING DATE: 2002-11-07

PRIOR APPLICATION NUMBER: PCT/IB02/00166

PRIOR PILING DATE: 2002-01-21

PRIOR APPLICATION NUMBER: US 10/050,902

PRIOR FILING DATE: 2002-01-18

PRIOR APPLICATION NUMBER: US 60/396,635

PRIOR APPLICATION NUMBER: US 60/331,045

PRIOR FILING DATE: 2001-11-07

NUMBER OF SEQ ID NOS: 170

PRIOR FILING DATE: 2001-11-07

NUMBER OF SEQ ID NOS: 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9, Application US/10622064 Publication No. US20040059094A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                     SEQ ID NO 9
LENGTH: 132
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LENGTH: 132
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                           Query Match
Best Local Similarity
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Best Local Similarity
                                                                         Matches
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APPLICANT: Maurer, Patrik F
APPLICANT: Maurer, Patrik F
ITTE OF INVENTION: Hapten-Carrier Conjugates and
FILE REFERENCE: 1700.0300001
CURRENT APPLICATION NUMBER: 101622,064
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,575
PRIOR FILING DATE: 2002-07-18
                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 33
SOFTWARE: PatentIn version 3.2
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APPLICANT: Spohn, Gunther
TITLE OF INVENTION: Antigen Arrays for Treatment of Bone Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                         FEATURE:
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                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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1 AKLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                                                    100.0%; Score 664; DB 15; nilarity 100.0%; Pred. No. 1.6e-66; Conservative 0; Mismatches 0;
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PRIOR FILING DATE: 2002-07-19
NUMBER OF SEQ ID NOS: 146
SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 20
                                                                                                                                                                                                                                                                                            Sequence 20, Application US/10622087
Publication No. US20040141984A1
GENERAL INFORMATION:
APPLICANT: Bachmann, Martin F
APPLICANT: Tissot, Alain
APPLICANT: Ortmann, Rainer
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NUMBER OF SEQ ID NOS: 93
SOFTWARE: PatentIn version 3.2
SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100
Best Local Similarity 100
Matches 132; Conservative
                                                           CURRENT APPLICATION NUMBER: US/10/622,087
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,639
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: US 60/470,432
PRIOR FILING DATE: 2003-05-15
                                                                                                                                                                                     APPLICANT: Frey, Peter
TITLE OF INVENTION: Amyloid Beta 1-6 Antigen Arrays
FILE REFERENCE: 1700.0350002
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APPLICANT: Fulurija, Alma
TITLE OF INVENTION: Ghrelin-Carrier Conjugates
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CURRENT FILING DATE: 2003-07-18
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TYPE: PRT
ORGANISM: Artificial Sequence
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Luond, Rainer
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Pred. No. 1.6e-66;
Mismatches 0;
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NUMBER OF SEQ ID NOS: 73
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 27
LENGTH: 132
TYPE: PRT
ORGANISM: Bacteriophage Q-beta
US-10-243-739-27
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US-10-243-739-27
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 Sequence 27, Application US/10244065
Publication No. US20030099668A1
GENERAL INFORMATION:
APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Maurer, Patrick
APPLICANT: Tissot, Alain
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GENERAL INFORMATION:
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Best Local 9
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Best Local (
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APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Lechner, Franziska
TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of
TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
FILE REFERENCE: 1700.0210001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
CURRENT FILING DATE: 2002-09-16
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Storni, Tazio
Maurer, Patrick
Tissot, Alain
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Pred. No. 3.5e-66;
1; Mismatches 0;
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Pred. No. 1.6e-66;
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APPLICANT: Meijerink, Edwin
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Cielens, Indulis
APPLICANT: Cielens, Indulis
APPLICANT: Renhofa, Regina
ITILE OF INVENTION: Method of Preparation and Use
ITILE OF INVENTION: Method of Preparation and Use
ITILE OF INVENTION NUMBER: US/10/244,065
CURRENT APPLICATION NUMBER: 0/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR FILING DATE: 2002-04-22
PRIOR FILING DATE: 2001-09-14
PRIOR FILING DATE: 2001-09-16
PRIOR FILING DA
                                                                                                                                          ; TYPE: PRT
; ORGANISM: Bacteriophage Q-beta-259
US-10-289-454-27
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Query Match
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Matches 131
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SEQ ID NO 27
LENGTH: 132
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PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/331,045
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CURRENT FILING DATE: 2003-02-10
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APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
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                                    Similarity
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       Conservative
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Score 661; DB 14;
Pred. No. 3.5e-66;
1; Mismatches 0;
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GENERAL INFORMATION:
APPLICANT: Renner, Wolfgang A.
APPLICANT: Hasch Alain
APPLICANT: Tissot, Alain
APPLICANT: Tissot, Alain
APPLICANT: Hesot, Alain
APPLICANT: Sebbel, Peter
APPLICANT: Lechner, Franziska
APPLICANT: Sebbel, Peter
TITLE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0190004
CURRENT APPLICATION NUMBER: US/10/050,902
CURRENT FILING DATE: 2001-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-9
PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/286,549
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-10-05
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 350
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 258
LENGTH: 132
TYPE: PRT
CORANISM: QD 259
US-10-050-902-258
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Search completed: January 4, 2005, 09:41:23 Job time : 33.7351 secs
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US-10-050-902-258
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Best Local Similarity 99.3
Matches 131; Conservative
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o. US20030175290A1
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99.2%; Pred. No. 3.5e-66;
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Matches 130; Conservative
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121 LIDAIDQLNPAY 132

61 NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDBERAFVRTBLAALLASPL 61 NYKVQVKIQNFTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL

120

60 60 0

A;Residues: 1-60 <sto> A;Residues: 1-60 <sto> R;Maita, T.; Konigsberg, W. J. Biol. Chem. 246, 5003-5024, 1971 A;Title: The amino acid sequence of the Qbeta coat protein. A;Reference number: A92088; MUID:71288580; PMID:5570434 A;Accession: A92088 A;Molecule type: protein A;Residues: 1-21, D', 23-55,57-132 <mai> C;Superfamily: phage GA coat protein Query Match 98.6%; Score 655; DB 1; Length 132; Best Local Similarity 98.5%; Pred. No. 1e-56; Matches 130; Conservative 1; Mismatches 1; Indels 0; Gaps 0;</mai></sto></sto>	A;Title: Determination of the first half of the coat protein cistron of bacteriophage Qbt A;Reference number: A92240; MUID:79048469; PMID:361741 A;Recession: A92240 A;Rolecule type: mRNA A;Residues: 1-80 <esc> A;Cross-references: UNIPROT:P03615 A;Cross-references: UNIPROT:P03615 R;Stoll, E.; Wilson, K.J.; Reiser, J.; Weissmann, C. J. Biol. Chem. 252, 990-993, 1977 A;Title: Revised amino acid sequence of Qbeta coat protein between positions 1 and 60. A;Reference number: A92221; MUID:77118576; PMID:838709 A;Rolecule type: protein</esc>	RESULT 1 VCBPQB coat protein - phage Q-beta C;Species: phage Q-beta C;Date: 29-Jul-1981 #sequence revision 24-Sep-1981 #text_change 09-Jul-2004 C;Accession: A92240; A92221; Ā92088; A04224 R;Escarmis, C.; Sastry, P.A.; Billeter, M.A. T Biol Chem 251 8390-8390, 1978	10.8 376 2 C84316 10.8 384 2 A45490 10.8 458 2 T49914 10.8 654 2 S69673 10.7 248 2 A86786 10.7 366 2 F70618 10.7 399 1 A4385 10.7 434 2 S72430 10.7 434 2 S00676 10.7 1461 2 A85547 ALIGNMENTS	72.5 10.9 129 72.5 10.9 129 72.5 10.9 129 72.5 10.9 129
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C;Accession: S22340
R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete se
A;Reference number: S22340; MUID:92182018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Dhaese, P.; Vandekerckhove, J.S.; Van Montagu, M.C. Eur. J. Biochem. 94, 375-386, 1979
A;Title: The primary structure of the coat protein of the A;Reference number: A04225; MUID:79148387; PMID:107028
A;Accession: A04225
A;Molecule type: protein
A;Residues: 1-131 < CHA>
A;Cross references: UNIPROT:P03616
C;Superfamily: phage GA coat protein
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C;Species: Listeria seeligeri
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C;Species: phage PRR1
C;Date: 30-Nov-1979 #sequence_revision 30-Nov-1979 #text_change
C;Accession: A04225
C;Accession: A04225
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A;Status: preliminary
A;Nolecule type: genomic RNA
A;Residues: 1-331 <INO>
A;Cross-references: UNIPROT:P09677; EMBL:X07489
A;Note: the authors translated the readthrough stopcodon TGA for residue C;Superfamily: phage GA coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Accession: S01964
R;Inokuchi, Y.; Jacobson, A.B.; Hirose, T.; Inayama, S.; Hira
Nucleic Acids Res. 16, 6205-6221, 1988
A;Title: Analysis of the complete nucleotide sequence of the
A;Reference number: S01963; MUID:88289362; PMID:3399390
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                                                                                            ;Date: 22-Nov-1993
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   B: complete sequence of ilo : MUID:92182018; PMID:1543752
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Pred. No. 0.00
14; Mismatches
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Pred. No. 3.3e-44;
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C;Genetics:
A;Gene: ATSP:F316.9
A;Map position: 1
A;Introns: 14/3; 281/3; 4
C;Superfamily: pyruvate c
                                c;Species: Listeria monocytogenes
A;Variety: strain 12067
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change
C;Date: 32-34231
R;Rasmussen, O.F.; Beck, T.; Olsen, J.E.; Dons, L.; Rossen, L.
Infect. Immun. 59, 3945-3951, 1991
A;Title: Listeria monocytogenes isolates can be classified into A;Reference number: 824230; MUID:92040062; PMID:1937753
A;Accession: S24231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cispecies: Arabidopsis thaliana (mouse-ear cress)
Cipate: 01-Feb-1999 #sequence_revision 01-Feb-1999 #text_change 09-Jul-2004
C;Accession: T00646
R;Federspiel, N.A.; Palm, C.J.; Conway, A.B.; Kurtz, D.B.; Conway, A.R.; Au, M.; A; Vysotekaia, V.S.; Yu, G.; Ecker, J.; Theologis, A.; Davis, R.W.
submitted to the EMBL Data Library, February 1998
A;Reference number: Z14197
A;Accession: T00646
A;Accession: T00646
A;Status: translated from GB/EMBL/DDBJ
A;Residues: 1-540 <FED>
A;Cross-references: UNIPROT:048683; EMBL:AC002396; NID:g2749918; PIDN:AAC00577.1;
A;Cross-references: UNIPROT:048683; EMBL:AC002396; NID:g2749918; PIDN:AAC00577.1;
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A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-530 <HAA>
A;Cross-references: UNIPROT:P31830; EMBL:X60462;
A;Note: the authors translated the codon GCC for C;Superfamily: dipeptide transport protein
                 A;Status: nucleic
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               acid sequence
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Nucleic Acids Res. 17, 6406, 1989
A,Title: Nucleotide sequence of the listeriolysin gene from a Listeria monocytogenes
A,Reference number: S05306; MUID:89366684; PMID:2505236
A,Accession: S05306
A,Accession: S05306
A,Molecule type: DNA
A,Residues: 1-529 < DOM>
A,Residues: 1-529 < DOM>
A,Cross-references: EMBL:X15127; NID:g44106; PIDN:CAA33223.1; PID:g44107
A,Experimental source: strain EGD
A,Cross-reference: derived from a weakly hemolytic strain, serotype 1/2a
A,Mote: this sequence is derived from a weakly hemolytic strain, serotype 1/2a
R,Mengaud, J.; Chenevett, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
Infect. Immun. 55, 3225-3227, 1987
A,Title: Identification of the structural gene encoding the SH-activated hemolysin of A,Accession: A47606; MUID:88057627; PMID:2824384
A,Accession: A47606
A,Status: preliminary
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A;Experimental source: strain 12067, serotype 4b
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1991
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Listeria monocytogenes
C;Species: Listeria monocytogenes
C;Date: 21-Oct-1992 #sequence revision 21-Oct-1992 #text_change 09-Jul-2004
C;Accession: A43505; S05306; A47606; S12400; A61079
C;Accession: A43505; S05306; A7606; S12400; A61079
R;Mengaud, J; Vicente, M.F.; Chenevert, J.; Pereira, J.M.; Geoffroy, C.; Gicquel-Sanzey
Infect. Immun. 56, 766-772, 1988
A;Title: Expression in Escherichia coli and sequence analysis of the listeriolysin O det
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C;Keywords: virulence factor
F;1-25/Domain: signal sequence #status predicted
A;Molecule type: DNA
A;Residues: 413-480 KMZ>
A;Residues: 43-480 KMZ>
A;Cross-references: GB:M29171
A;Cross-references: GB:M29171
A;Cross-references: GB:M29171
A;Cross-references: GB:M29171
A;Michel, E.; Reich, K.A.; Favier, R.; Berche, P.; Cossart, P.
Mol. Microbiol. 4, 2167-2178, 1990
A;Title: Attenuated mutants of the intracellular bacterium Listeria
A;Reference number: S12400; MUID:91211627; PMID:1965218
A;Recession: S12400
A;Molecule type: DNA
A;Residues: 483-493 KMIC>
A;Experimental source: strain LO28, serotype 1/2c
C;Genetics:
A;Gene: hlyA; lisA
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A;Note: this sequence is derived from a strongly hemolytic strain, serotype 1/2c
R;Domann, E; Chakraborty, T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Reference number: A43505;
A; Accession: A43505
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A; Residues: 1-529 < MEN>
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D.; Jones, L.M.; Karst, U.
Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; Maitournam, ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; Voss, H.; We A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Accession: AC1100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Superfamily: dipeptide transport protein C;Keywords: virilence factor F;1-25/Domain: signal sequence #status predicted <NIG>F;26-529/Product: listeriolysin O #status predicted <NI
                                               ivanolysin precursor - Listeria ivanovii
(;Species: Listeria ivanovii
C;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 09-Jul-2004
C;Accession: $22341; $36683
C;Accession: $22341; $36683
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C;Date: 27-Nov-2001
C;Accession: AC1100
C;Accession: S22341; S36683
R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete sequence of ilo: A;Reference number: S22340; MUID:92182018; PMID:1543752
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A; Residues: 1-529 <G
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C;Superfami
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Pred. No. 2
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Wehland,
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Cell 56, 291-301, 1989
A;Title: Functional dissection and sequence of yeast HAP1 activator.
A;Reference number: A31312; MUID:89106221; PMID:2643482
A;Reference number: A31312; MUID:89106221; PMID:2643482
A;Residues: 1-144,'I',146-322,'R',324-454,'N',456-507,'M',509-586,'K',588-882,'N',884-95
A;Cross-references: EMBL:JU03152; NID:g171645; PIDN:AAA34662.1; PID:g171646
A;Cross-references: EMBL:JU03152; NID:g171645; PIDN:AAA34662.1; PID:g171646
A;Cross-references: EMBL:JU03152; NID:g171645; P.P.
J. Mol. Biol. 204, 263-276, 1988
A;Title: CYP1 (HAP1) regulator of oxygen-dependent gene expression in yeast. I. Overall A;Reference number: S15447; MUID:89125585; PMID:2851658
A;Accession: S15447
A;Rossidues: 1-62,'R',64-1305,'Y',1306-1470,'LVDFYRADFPIWE' <CRE1>
A;Cross-references: EMBL:X1305,'Y',1306-1470,'LVDFYRADFPIWE' <CRE1>
A;Cross-references: EMBL:X1305,'Y',1306-1470,'LVDFYRADFPIWE', <CRE1>
A;Cross-references: EMBL:X1305,'Y',1306-1470,'LVDFYRADFPIWE',1305,'YDFYRADFPIWE',1305,'YDFYRADFPIWE',1305,'YDFYRADFPIWE',
C;Genetics:
A;Gene: SGD:HAP1; CYP1; MIPS:YLR256w
A;Cross-references: SGD:S0004246; MIPS:YLR256w
A;Map position: 12R
C;Superfamily: regulatory protein HAP1; GAL4 z:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Molecule type: DNA
A;Residues: 1-1502 <00H>
A;Residues: 1-1502 <00H>
A;Cross-references: EMBL:U20865; NID:g662330; PIDN:AAB67387.1; PID:g662331; GSPDB:GN0001
A;Experimental source: strain S288C (AB972)
R;Pfeifer, K.; Kim, K.S.; Kogan, S.; Guarente, L.
Cell 56, 291-301, 1989
Cell 56, 291-301, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RiJohnson, D.

RiJohnson, D.

Submitted to the EMBL Data Library, February submitted to the EMBL Data Library, February
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NALternate names: protein 19672.1; protein YLR256w; regulatory protein CYP.
C;Species: Saccharomyces cerevisiae
C;Date: 30-Sep-1991 #sequence revision 23-Feb-1996 #text_change 12-Nov-1999
C;Accession: S59400; A31312; S15447; S05804; S15446
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C;Superfamily: dipeptide transport protein
F;1-24/Domain: signal sequence #status predicted
F;25-528/Product: ivanolysin #status predicted <M
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A; Residues: 1-319, 'T', 321-528 < KRE
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A; Molecule type: DN
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A; Residues: 1-528 < HAS>
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Best Local (
                                                                                                                                                                                     Note: the sequence is from mutant CYP1-18
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Pred. No. 5
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hypothetical protein COS41.6 - sea squirt (Ciona intestinalis) C;Species: Ciona intestinalis C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 09-Jul-2004 C;Accession: T31660 R;Bird, A.P.; Clark, V.; Jones, S.J.; Leitgeb, S.; Dobson, R.; Tweedie, S. submitted to the EMBL Data Library, December 1996
                                                                                                                                     T31660
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C; Superfamily:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-282 < KUR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Title: Genome sequence of Yersinia pestis, the causative A;Reference number: AB0001; MUID:21470413; PMID:11586360 A;Accession: AI0186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probable iron-siderophore
C;Species: Yersinia pestis
C;Date: 02-Nov-2001 #seque
C;Accession: AI0186
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F;245-445/Domain: heme binding thathus predicted <HEM>
F;245-445/Domain: heme binding 3-39-394,415-420/Region:
F;299-304,323-328,347-352,373-378,389-394,415-420/Region:
F;1308-1481/Domain: activation element #status predicted
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                                                                                                                                                         RESULT
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                                                                                                                                                                                                                                                                    127
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                                                                                                                                                                                                                                                                                                              103 LLSQGASISETLTVFDLVSRGRYAHQSF
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                                                                                                                                                                                                                                                                                                                                                                                                                                         13 KDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVKIQNPT
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                                                                                                                                                                                                                                                                                                                                                       73 ACTANGSCDPSVT-----RQKYADVTFSFTQYSTDEERAFVRTELAALLASPLLIDAID 126
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                                                                                                                                                                                                                                                                    QLN 129
                                                                                                                                                                                                                                                                                                                                                                                                KNGAFSVIIGPNGCGKSTLLRALSRS-----LLPQNGSIRLDQ--QDIQHYKAKVFARQLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SVTRQKYADV-TFSFTQYSTD-EERAFVRTELAALLASP-LLIDAIDQLN 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRGISPKPSNGLSSVQPLLSSFSMNQLNGGTIPTVPSLTNITSQMGALPSLDRITTNQIN 1337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inner membrane protein malk; ATP-binding cassette homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12.1%;
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22.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      37; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 79;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 80.5;
Pred. No. 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GB:AL590842; PIDN:CAC90356.1; PID:g15979576; GSPDB:GN
                                                                                                                                                                                                                                                                                                            - FHQWSTEDER-IVKAAISAVNLESLVQQRVS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATP-binding component YPO1533 [imported]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 2; Length 282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ----SQAGAVPALEKRVTVSVS
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S.; Barrell,
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A;Cross-references: UNIPROT:P40850; EMBL:X89016; NID:g887621; PID:g887626 R;Vermut, M.; Widner, W.R.; Dinman, J.D.; Wickner, R.B. Yeast 10, 1477-1479, 1994 A;Title: Sequence of MKTI, needed for propagation of M(2) satellite dsRNA A;Reference number: S50279; MUID:95176705; PMID:7532890 A;Accession: S50279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Saccharomyces cerevisiae
C;Species: 10-Oct-1995 #sequence revision 10-Nov-1995 #text_change 09-Jul-2004
C;Accession: S57537; S50279; S63024; S63017; S65096
R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M. submitted to the EMBL Data Library, June 1995
                                                                                                                                                                                                                  A;Experimental source: strain S288C

R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.;

Yeast 12, 485-491, 1996

A;Title: The sequence of a 17 933 bp segment of

A;Reference number: S65092; MUID:96310628; PMID:
                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Poehlmann, R.; Philippsen, P. submitted to the Protein Sequence Database, A;Reference number: S62997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: DNA
A;Molecule: 1-830 <SOW>
A;Cross-references: EMBL:Z71361; NID:g1301982; PID:g1301983; MIPS:YNL085w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-29,'G',31-808,'TMKTCLNYH' <VER>
A;Residues: 1-29,'G',31-808,'TMKTCLNYH' <VER>
A;Cross-references: EMBL:U09119; NID:9520475; PIDN:AAC49470.1;
R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M.
submitted to the Protein Sequence Database, April 1996
A;Reference number: S63018
A;Accession: S63024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Reference number: S57533
A; Accession: S57537
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A; Residues: 1-432 <BIR>
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                                                                                                                                                                                             A; Reference number: S65092;
A; Accession: S65096
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A;Cross-references:
A;Map position: 14L
                                                                                                                       A;Molecule type: DNA
A;Residues: 1-830 <SC
                                                 A;Gene: SGD:MKT1
                                                                                                                                                                                                                                                                                                                                         ;Molecule type: DNA
;Residues: 569-830 <POE>
;Cross-references: EMBL;Z71361; MIPS:YNL085w
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                                                                                                  Cross-references:
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                                                                                                                    1-830 <SOF>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DTSSEGEVTKDGPGNLAENP---TPSNAARBLQESVASEVLETTVVKSAIQ-EQDSSAYR
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mes: protein N2302; protein YNL085w
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                                                                                                                                                                        acid sequence not shown
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                         SGD:S0005029; MIPS:YNL085w
                                                                                             EMBL:X89016; NID:g887621; PIDN:CAA61425.1; PID:g887626
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Pred. No.
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R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, L. erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClella ; Karp, P.; Romero, P.; Zhang, S. Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm, F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Superfamily: Saccharomyces cerevisiae MKT1 protein C;Keywords: transmembrane protein F;615-631/Domain: transmembrane #status predicted <TVM/
                                                          A;Molecule type: DNA
A;Residues: 1-136 <KUR>
A;Cross-references: UNIPROT:Q8U8F3; GB:AE008689; PIDN:AAL44939.1; PID:g17742593; GSPDB:G
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A; Map position: li
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Science 294, 2323-2328, 2001
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                                                                                                                                                                  A; Reference number: A; Accession: AB3065
                                                                                                                                                                                      A;Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58 A;Reference number: AB2577; MUID:21608550; PMID:11743193
                                                                                                                                                                                                                                                                                                                                                                                                                                  conserved hypothetical protein Atu4139 [imported] -
C;Species: Agrobacterium tumefaciens
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A;Reference number: A97359; MUID:21608551; PMID:11743194
A;Gene:
                                             A;Experimental source: strain
                                                                                                                                                                                                                                                                                                                                                                                  C; Accession: AE3065
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C.; Allinger, M.; Doughty, D.; Scott, C.; Lappas,
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C58, Cerec

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PubMed=14667253;

Bacher J.M., Bull J.J., Ellington A.D.;

Bacher J.M., Bull but chemically ambiguous proteomes.";

"Evolution of phage with chemically ambiguous proteomes.";

BMC Evol. Biol. 3:24-24(2003).

EMBL; AY099114; AAM33126.1; -.
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EMBL; V00643; CAA23992.1; -.
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license@isb-sib.ch).
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01-OCT-2002 (TrEMBLre1. 2
05-JUL-2004 (TrEMBLre1. 2
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AAA16663;
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EMBL; AY099139; AAA16663.1; -.
HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; IEA.
InterPro; IPR002703; Levi_coat.
InterPro; IPR000504; RNA rec mot.
Pfam; PF01819; Levi_coat; 1.
PROSITE; PS00030; REM RNP 1; UNKNOWN 1.
SEQUENCE 329 AA; 36134 MW; E0F30215A8C197CB CRC64;
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PubMed=14667253;
Bacher J.M., Bull J.J., Ellington A.
"Evolution of phage with chemically
BMC Evol. Biol. 3:24-24(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Al read-through protein (Al protein).
Bacteriophage Q-beta.
Viruses; sBRNA positive-strand viruses,
Allolevivirus.
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MEDLINE=94109687; PubMed=7506687;
Kozlovska T.M., Cielens I., Dreilinna D.,
Ose V., Pumpens P.;
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Pred. No. 4.8e-55;
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01-MAY-2000
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MEDLINE=94109687; PubMed=7506687;
KOzlovska T.M., Cielens I., Dreilinna
Ose V., Pumpens P.;
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Viruses; ssRNA positive-strand
Allolevivirus.
            Pfam; PF01819; Levi_coat; Coat protein
                             Beekwilder M.J., Nieuwenhuizen R., van Duin J.;
Submitted (ApR-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF05924; AAC14699.1; -
HSSP; P03615; 1QBE.
GO; GO:0005198; C:viral capsid; IEA.
GO; GO:0005198; C:viral capsid; IEA.
GO; GO:005198; C:viral capsid; IEA.
Interpro; IPR002703; Levi_coat.
                                                                                                                           SEQUENCE FROM N.A.

MEDIJINE=9619948; PubMed=8609616;

Beekwilder J., Nieuwenhuizen R., Poot R., van Duin J.;

Beekwilder structure model for the first three domains

Control of A-protein synthesis.";

J. Mol. Biol. 256:8-19(1996).
                                                                                                                                                                                                                                                                                              Major coat protein.
Enterobacteria phage MX1.
Viruses; ssRNA positive-strand
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EMBL; M99039; AAA16663.1; -.
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02-MAR-2004
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01-JUN-2003
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AP059242; ACC14700.1; -.

R HSSP; P03615; 1QBE.

GO; GO:0019028; C:viral capsid; IEA.

GO; GO:0005198; F:structural molecule activity; IEA.

InterPro; IPR002703; Levi_coat.

InterPro; IPR002703; Levi_coat.

InterPro; IPR000504; RNA_rec_mot.

R Ffam; PF01819; Levi_coat; 1.

R PROSITE; PS00030; RRM RNP 1; UNKNOWN 1.

SEQUENCE 329 AA; 35880 MW; 372B1DD5DDB52F15 CRC64;
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A1-protein.
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"Secondary structure model for the
Control of A-protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
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Pred. No. 2e-46;
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Matches 106
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01-AUG-1998 (
01-AUG-1998 (
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Q9T0S0;
01-MAY-2000
                                                                                  Bacteriophage M11.
Viruses; ssRNA positive-strand
Allolevivirus.
    SEQUENCE FROM N.A. MEDLINE=95239761;
                                                              NCBI_TaxID=74336;
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Mol. Biol. 247:9
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GO:0005198; F:structural molecule
erPro; IPR002703; Levi coat.
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  PubMed=7723040;
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80.3%; Pred. No. 6.1e
tive 12; Mismatches
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P09673;
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SEQUENCE FROM N.A.
MEDLINE-88289362; PubMed=3399390;
MITASHIMA A., Hirose T., Inayama S.,
Mirashima A., momplete nucleotide
  EMBL;
                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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"Analysis of the complete coliphage SP.";
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GO; GO:0005198; F:structural molecule activity;
InterPro; IPR002703; Levi coat.
InterPro; IPR000504; RNA_rec_mot.
Pfam; PF01819; Levi coat; 1.
PROSITE; PS00030; RRM_RNP_1; UNKNOWN 1.
SEQUENCE 329 AA; 35893 MW; 3E33CD821EE625F4
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Allolevivirus.
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HSSP; P03615; 1QBE.
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MEDLINE=96190948; PubMed=8609616;
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FUNCTION:
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RESULT 10
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                                                                                               Best Local Similarity Matches 104; Conserv
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                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for comentities requires a license agreement (See http://www.isb-sib.ch/an
                                                                                                                                                                                                                                                                                                                                                                                                     coliphage SP.";
Nucleic Acids Res. 16:6205-6221(1988).
-!- FUNCTION: Not yet known.
-!- MISCELLANEOUS: The readthrough protein Al includes the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteriophage SP.
Viruses; ssRNA positive-strand viruses, no DNA stage; Allolevivirus.
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01-MAR-1989 (Rel. 10, Last sequence update)
29-MAR-2004 (Rel. 43, Last annotation updat
Readthrough protein Al [Contains: Coat prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
Coat protein; RNA-binding.
SEQUENCE 132 AA; 14129 MW;
                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                  HSSP; P03615; 1QBE.
InterPro; IPR002703; Levi_co
Pfam; PF01819; Levi_coat; 1.
                                                                                                                                                                                                                                            EMBL; X07489; CAB37299.1; PIR; S01964; S01964.
                                                                                                                                                                                                                                                                                     or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hirashima A., Hirose T., Inayama S., Inokuchi "Analysis of the complete nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=88289362; PubMed=3399390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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NYKVQVKIQNETACTANGSCDESVTRQKYADVIESFTQYSTDEERAFVRTELAALLASPL
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                                                            AKLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                           AKLNQVTLSKIGKNGDQTLTLTPRGVNPTNGVASLSEAGAVPALEKRVTVSVAQPSRNRK
                                                                                                                                                   331 AA;
                                                                                               Conservative
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                                                                                             Score 531.5; DB 1
Pred. No. 4.5e-43;
2; Mismatches 15
                                                                                                                                                Coat protein.
Readthrough protein Al.
, 4A6642E4B52C6582 CRC64;
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Pred. No. 1.5e-43;
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Best Local
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01-AUG-1998
01-MAR-2004
Al-protein.
Al-protein.
Enterobacteria phage NL95.
Enterobacteria positive-strand viruses,
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HSSP; P036.15; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity;
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
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01-MAY-2000
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"Secondary structure model for the first three
Control of A-protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
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MEDLINE=96190948; PubMed=8609616;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Secondary structure model RNA phage Q beta.";
J. Mol. Biol. 247:903-917(1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beekwilder M.J., Nieuwenhuizen R., "Secondary structure model for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. MEDLINE=95239761; PubMed=7723040;
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Allolevivirus.
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                                                                                                    (TrEMBLrel.) (TrEMBLrel.)
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                                      no DNA stage; Leviviridae;
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NCBI_TaxID=75725; Allolevivirus.

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COAT BPPRR
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DT 21-JU
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                                                                                                                       P03616;
21-JUL-1986
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COAT_BPPRR
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GO; GO:0005198; F:structural molecule activity;
InterPro; IPR002703; Levi coat.
InterPro; IPR002703; Nar rec_mot.
Pfam; PF01819; Levi coat; 1.
PROSITE; PS00030; RRM_RNP_1; UNKNOWN 1.
SEQUENCE 330 AA; 36175 NW; 961E55F408334410
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J. Mol. Biol. 247:903-917(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Viruses; ssRNA positive-strand viruses, unclassified Leviviridae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Coat protein.
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05-JUL-2004
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Submitted (APR-1998) to the EME
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   SEQUENCE
                           InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
Coat protein; Direct protein se
                                                                                                                                                                                                                                  bacteriophage PRR1."
                                                                                                                                                                                                                                                         The primary structure of the
                                                                                                                                                                                                                                                                                                                              MEDLINE=79148387; PubMed=107028;
                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=12024;
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      131 AA;
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(Rel. 01,
(Rel. 44,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
   protein sequencing; RNA-binding.
14535 MW; E7E639E1E50FC612 CRC64;
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coat protein of
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01-JUL-1993 (Rel. 26, Creat
01-JUL-1993 (Rel. 26, Last
05-JUL-2004 (Rel. 44, Last
Seeligeriolysin precursor (
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Q8VDC2;
01-MAR-2002
01-MAR-2002
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Mus musculus (Mouse).
Thervota; Metazoa; Chordata;
Therval; Rodentia;
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Mamm. Genome 13:646-655(2002).
EMBL; AJ428064; CAD20986.1; -.
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MEDLINE=22350095; PubMed=12461651;
Kiss H., Darai E., Kiss C., Kost-A
                                         Name=lso;
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                     Listeria seeligeri.
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GO:0016021; C:integral to membrane;
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                                                                                                                                                                                                                                                                                                                                                                                                     47 RVTVSVSQPSRNRKNYKVQV------KIQNPT-----ACTANGSCDPSVTRQ-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 OTLVLNPRGVNPT------NGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   473 AA;
                                                        (Rel. 26, Created)
(Rel. 26, Last sequence update)
(Rel. 44, Last annotation update)
ysin precursor (Thiol-activated cy
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Pred. No. 0.0028
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Sciurognathi;
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Bacteria;

Firmicutes;

Bacillales;

Listeriaceae;

Listeria

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Search completed: January 4, 2005, 09:16:17 Job time: 37.2676 secs
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Best Local Similarity
Matches 34; Conserv
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SIGNAL
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HEBLINE-92182018; MEDINE-92182018; MEDINE-921820182018; MEDINE-92182018; MEDINE-921820182018; MEDINE-92182018; MEDINE-92182018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; X60462; CAA42996.1; -.
HSSP; P19995; 1PFO.
InterPro; IPR001869; Thiol cytolysin.
Pfam; PF01289; Thiol cytolysin; 1.
PRINTS; PR01400; TACTTOLYSIN;
PRODOM; PD077062; Thiol cytolysin; 1.
PROSITE; PS00481; THIOL CYTOLYSINS; 1.
CYTOLYSIS; Hemolysis; Lipid-binding; Signal; Toxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                            209
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                                                                                                                                                                                                                                                                                                                    49 TVSVSQPSRNRKNYKVQVKIQNPTACTANGSCDPSVTR--QKY------ADVTFSFTQ 98
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485
530 AA;
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485 Bi
; 59181 MW;
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25.0%; Pred. No. 5.2;
vative 22; Mismatches
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Binding to cholesterol (By similarity).
416F7A4DD2029866 CRC64;
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GenCore version (c) 1993 - 2005

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ABG94233	ADK52196	ADJ67162	ADJ36316	ADK17147	ADJ82059	ADD24133	ABR44558	ABU09693	ABR56455	ABG80632	ABG94320	ADK52197	ADJ67163	ADJ36317	ADK17148	ADJ82060	ADD24134	ABR44559	ADOCOOCA.
Abg94233	Adk52196	Adj67162	Adj36316	Adk17147	Adj82059	Add24133	Abr44558	Abu09693	Abr56455	Abg80632	Abg94320	Adk52197	Adj67163	Adj36317	Adk17148	Adj82060	Add24134	Abr44559	ACOCOURAL PROPERTY AND ACCOURAGE AND ACCOURACE AND ACCOURAGE AND ACCOURA
Bacteriop	Bacteriop	Phage Qbe	Bacteriop	Virus-lik	Protein	Bacteriop	Bacteriop	Bacteriop	Bacteriop	Bacteriop	pQB251 pr	Bacteriop	Phage Qbe	Bacteriop	Virus-lik	Protein	Bacteriop	Bacteriop	JOHNSON

ALIGNMENTS

ABG94318 standard;

protein;

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10-DEC-2002
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19-JAN-2001; 2001US-0262379P.
04-MAY-2001; 2001US-0288549P.
05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-0331045P.

(CYTO-) CYTOS BIOTECHNOLOGY AG.

Renner WA, Piossek C; Bachmann M, Tissot Þ Maurer η, Lechner Ţ Sebbel

P,

WPI; 2002-627351/67.

diseases. Molecular antigen array used in the production of vaccines for infectious

Claim 18; Page 145; 441pp; English.

This invention relates to a novel ordered and repetitive antigen array used in the production of vaccines for infectious diseases. The invention also discloses a composition comprising a non-natural molecular scaffold comprising a core particle selected from a core particle of a non-natural origin and a core particle of natural origin and an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond. Also disclosed is an antigen or antigenic determinant with at least one second attachment

where the antigen or antigenic determinant is amyloid beta peptide

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 132;
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Best Local :
                                           04-MAY-2001;
05-OCT-2001;
07-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Molecular antigen array; vaccine; antigen; antimicrobial; molecular scaffold; amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgE-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus erythematosus; osteoporosis; inflammatory immune disease; pupus erythematosus; multiple sclerosis; inflammatory immune disease; lymphadenopathy; Alzheimer's disease; angioimmunoproliferative disease; lymphadenopathy; Alzheimer's disease; angioimmunoproliferative lymphadenopathy; immunoblastive lymphadenopathy; rheumatoid arthritis; diabetes; infectious disease; factor Xa;
                                                                                                                                                                         19-JAN-2001;
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                                           ; 2001US-0262379P.
; 2001US-0288549P.
; 2001US-0326998P.
; 2001US-0331045P.
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CC molecular Scaffold comprising: (1) a core particle selected from: (1) a core particle of a non-natural origin; and (2) a core particle of natural corigin; and (2) a core particle by at least cone covalent bond; (b) an antigen or antigenic determinant with at least one second attachment site, where the antigen or antigen or antigenic determinant is camploid beta peptide (Abeta 1-42) or its fragment, and where the second courring with the antigen or antigenic determinant; and (ii) an cocurring with the antigen or antigenic determinant; and (ii) an cocurring with the antigen or antigenic and cocurring with the antigen or antigenic cocurring with the antigen or antigenic action cocurring with the antigen array. The composition is used in consumation and as a vaccine for disease such as influenza, graft corrected and repetitive antigen array. The composition is used in courring with the antigen array influence and antigenic cocurring and influences. The composition is used in court lympholastic leukaemia, non-Hodgkin's lymphoma, Grave's disease, systemic lympholastic particle with disease, allergic asthma, cocurring with the antigen array influences and infectious diseases. The antigens are modified to possess a cleavage site (enterokinase or factor Xa) and a Cysteine- containing N-coriginal linker peptide which serves as the attachment point to a cyrus like particle or bacterial protein or peptide which 
Sequence 132 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a composition comprising: (a) a non-natural molecular scaffold comprising: (i) a core particle selected from: (1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 18; Page 145; 418pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Molecular antigen array used in the production of vaccines for infectious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-636514/68.
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(NOVS)
(MAUR/)
(LECH/)
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ORTMANN R.
LUEOEND R.
STAUFENBIEL M.
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Bachmann M,
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, Tissot A,
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Sebbel
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RESULT 3
ABR56454
ID ABR5
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Best Local Similarity
Matches 132; Conserv
 ABR56454 standard;
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                                                                                                          LIDAIDQLNPAY 132
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                                                                                                                                                                                                                                                                        Conservative
 protein;
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Pred. No. 8.2e-71
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Length 132;

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Gaps

120 60 60

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132

cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory; antithyroid; antidiabetic; neuroprotective; notropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis; Alzheimer's disease; osteoporosis; rheumatoid arthritis;

Bacteriophage Obeta.

inflammatory autoimmune disease.

Bacteriophage Q-beta coat protein mutant SEQ ID NO:25

28-JUL-2003 ABR56454;

(first entry)

Antigen presenting cell; APC; immune response; virus like particle;

VLP;

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The present invention describes a composition (C) for enhancing an immune CC response against an antigen or a virus-like particle (VLP) bound to at least one antigen, or a CC comprises a virus-like particle (VLP) bound to at least one antigen, or a CC VLP capable of being recognised by the immune system of the animal. Also described: (1) enhancing an immune response against an antigen or a VLP cin an animal comprising introducing (C) into the animal; (2) vaccines CC comprising (C) together with a pharmaceutical diluent, carrier or CC excipient; (3) immunising or treating an animal comprising administering the vaccine; and (4) enhancing anti-viral CC the animal by administering the vaccine; and (4) enhancing anti-viral CC canimal by administering the vaccine; and (4) enhancing anti-viral CC canimal comprising introducing (C) into the animal. (C) CC antislergic, immunosuppressive, antiaddictive, antiinflammatory, CC antishlergic, immunosuppressive, antiaddictive, antiinflammatory, contractive, antiinflammatory, contractive, and antiarthritic activities. (C) or the vaccines can be antishred to an immune response against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a VLD in an insurance against an antigon or a vLD in an insurance against an antigon or a vLD in an insur
                                    Query Match
Best Local (
Matches 132;
                                                                                                                                                                                                                                                                                                                                        used for enhancing an immune response against an antigen or a VLP in an animal, enhancing anti-viral protection in an animal, or immunising or treating tumours and infectious diseases such as viral, bacterial, parasitic or fungal infections. The vaccine compositions are also useful for preventing or treating allergies, drug addiction, graft-versus-host disease, Crohn's disease, Grave's disease, diabetes, multiple sclerosis, allergies of the series of the serie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A composition, useful for enhancing an immune response against an antigen or a virus-like particle, enhancing anti-viral protection in an animal, or immunizing or treating tumors or infectious diseases, e.g. viral
                                                                                                                                                          Sequence 132
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                                                                                                                                                                                                                                                                               autoimmune disease.
                                                                                                                                                                                                                                                                                                                Alzheimer's disease, osteoporosis,
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                                        Local Similarity
    Conservative
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                                                                                                                                                                                                                       , osteoporosis, rheumatoid arthritis, or inflammatory ACC69838 to ACC69852 and ABR56401 to ABR56509 used in the exemplification of the present invention
                                        100.0%;
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                                664; DB 6;
No. 8.2e-71;
                                                                         Length 132;
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1 ARLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60

Query Match Best Local Similarity

100.0%;

Score 664; DB 6; Pred. No. 8.2e-71;

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                                          The invention describes an angiotensin peptide moiety carrier conjugate comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with remin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kidney failure, and retina haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta cost protein used in the preparation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-0331045P;
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-IB000166.
19-JUL-2002; 2002US-0396637P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteriophage Qbeta; coat protein; hypotensive; cerebroprotective; cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction; congestive heart failure; kidney failure; retinal haemorrhage; mutant;
                                                                                                                                                                                                                                                                                                          New angiotensin peptide moiety carrier conjugate comprising a carrier and an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension
                                                                                                                                                                                                                                                           Claim 20; Page 94; 97pp; English.
                                                                                                                                                                                                                                                                                            or infarction.
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                                                                                                                                                                                                                                                                                                                                                                                                                   Bachmann
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                                                                                                       and retinal
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immunostimulatory substance. The immunostimulatory substance is bound the VLP. Also described: (1) enhancing an immune response in an animal introducing (C) into the animal; (2) producing (C) for enhancing an immune response in an animal; (3) vaccines comprising (C) together with pharmaceutical diluent, carrier or excipient; and (4) immunising or
                                                                                                                                                                                                                                                                                      New compositions comprising immunostimulatory substances packaged into virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, tumors viral infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maurer P,
Pumpens P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunostimulatory; virus-like particle; bacteriophage; HBV; LCMV; hepatitis B virus; lymphocytic choriomeningitis virus; vaccine; immunostimulant; cytostatic; accidence; virucide; antibacterial; immune response; immunisation; allergy; tumour; breast cancer; neuroblastoma; leukaemia; viral disease; influenza; hepatitis; measles; chicken pox; bacterial infection; tuberculosis; pneumonia; syphilis.
                                                                                                                                                 The present invention describes a composition (C) for enhancing an immune response in an animal (C) comprises a virus-like particle (VLP), and an
                                                                                                                                                                                                                                Disclosure; Page 260-261; 322pp;
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CIELENS I.
RENHOFA R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SCHWARZ K.
MEIJERINK E.
LIPOWSKY G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MAURER P.
TISSOT A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CYTOS BIOTECHNOLOGY AG
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Cielens 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schwarz K, |
, Renhofa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  132 AA.
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                                                                                                                                                                                                                                  English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Meijerink E,
, Bachmann MF,
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F, Storni
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Best Local S
Matches 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vaccine composition; virus-like particle; core particle; first attachment site; antigen; antigenic determinant; prion PrP; PrP peptide; vaccine; neuroprotective; antiinflammatory; prion disease; Bovine Spongiform Encephalopathy; BSE;
                                                                             WPI; 2003-598483/56.
                                                                                                                                                                                                                                                                                                  18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-IB000166.
08-JUL-2002; 2002US-0393725P.
18-JUL-2002; 2002US-0396590P.
                                                                                                                                                   Bachmann M,
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Pred. No. 8.2e-71;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 132; Conserv
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like or a core particle with at least one first attachment site and at
least one antigen or antigenic determinant that is a prion protein (PrP)
or its dimer, or a PrP paptide, the antigen or antigenic determinant
being bound to the virus-like or core particle. The vaccine of the
invention may have neuroprotective or antinflammatory activity. The
composition is useful as a medicament or in manufacturing a medicament
for the treatment or prevention of prion diseases. The prion diseases may
include Bovine Spongiform Encephalopathy (BSE) or Creutzfeldt-Jakob
Disease. The present sequence is the amino acid sequence of a mutant coat
protein from bacteriophage Obeta which may be used during the creation of
the vaccine composition of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    phage) an particle.
                                                                                                                                                                                                                                                                                                       07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00059902.
21-JAN-2002; 2002WO-IB000166.
19-JUL-2002; 2002US-0396635P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Creutzfeldt-Jakob Disease) comprises a virus-like particle (e.g. RNA-phage) and at least one prion protein or peptide bound to the virus-like
New compositions comprising a core particle and at least one antigen antigenic determinant, useful for as a vaccine for therapy or prophylof bone diseases, particularly mammalian encephalopathies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-NOV-2002; 2002WO-EP012449.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003039225-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                osteopathic; vaccine; core particle; antigenic determinant; RANKL; bone disease; encephalopathy; immune system stimulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-MAY-2004
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                                                                                                                          WPI; 2003-441430/41.
                                                                                                                                                                                                                                           (CYTO-)
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                                                                                                                                                                                                                                              CYTOS BIOTECHNOLOGY AG
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                                                                                                                                                                                   Maurer
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Pred. No. 8.2e-71;
0; Mismatches 0;
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Matches 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antiallergic; antiasthmatic; cytostatic; vaccine; virus-like particle; interleukin; IL-5; IL-13; eotaxin; repetitive antigen array; allergic eosinophilic disease; asthma; Hodgkin's lymphoma.
                                                                                                                                                                                                                                                                              07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00059902.
21-JAN-2002; 2002WO-IB000166.
19-JUL-2002; 2002US-0396636P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                       07-NOV-2002; 2002WO-EP012455
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Pred. No. 8.2e-71;
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Composition comprising an ordered and repetitive antigen or antigenic determinant array, useful as a medicament, or for manufacturing a medicament for treating allergic eosinophilic diseases, e.g. asthma,

Hodgkin's lymphoma

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RESULT 9
ADJ36315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 132; Conserv
New compositions comprising a virus-like particle (VLP), an immunostimulatory substance bound to the VLP, and an antigen mixed the VLP, useful for enhancing immune response or for treating e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immune response; virus-like particle; immunostimulatory; allergy;
chronic disease; chronic viral disease; bacteriophage Qbeta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 132
                                                                                                                                    WPI; 2004-108361/11
                                                                                                                                                                                                  Bachman
                                                                                                                                                                                                                                                                                                                         20-JUN-2002; 2002US-0389898P
                                                                                                                                                                                                                                                                                                                                                                                        20-JUN-2003; 2003WO-EP006541
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage Qbeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antiallergic; cytostatic; virucide; immunostimulant; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteriophage Qbeta coat protein virus-like particle mutant N10K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ36315 standard; protein;
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                                                                                                                                                                                                                                                              (CYTO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; VLP; adjuvant; mutant; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    121 LIDAIDQLNPAY 132
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                                                                                                                                                                                                  MF,
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                                                                                                                                                                                               Renner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          245pp;
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Best Local Similarity 100.
Matches 132; Conservative
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antigenic a
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The invention relates to a new composition comprising: (i) a core particle with at least one first attachment site; and (ii) at least one antigen or antigenic determinant with at least one second attachment
                                                                                                                                                                New composition comprising a core particle having a first attachment and an antigen or antigenic determinant which is a ghrelin or ghrelin peptide having a second attachment site, useful for treating obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADJ67161 standard;
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                                                                                                            Disclosure; SEQ ID NO 19; 175pp; English
                                                                                                                                                                                                                                                                                                                                     Bachmann MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUL-2002; 2002US-0396638P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JAN-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADJ67161;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-JUL-2003; 2003WO-EP007849
                                                                                                                                                                                                                                                                                  2004-132866/13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; core particle; array.
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Pred. No. 8.2e-71;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               determinant; ghrelin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 132;
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                                                                                                                                                                     or ghrelin
ng obesity.
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Query Match
Best Local Similarity
Matches 132; Conserv
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                                                                                                                                                            19-JUL-2002; 2002US-0396639P.
15-MAY-2003; 2003US-0470432P.
                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                      neuroprotective; nootropic; vaccine; amyloid beta 1-6 peptide; core particle; antigen array; Alzheimer's disease; RNA bacteriophage; coat protein; CP; mutant; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteriophage Qbeta coat protein mutant K2R K13R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK52195
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 132 AA;
                                WPI; 2004-203731/19
                                                                                                                                                                                                           18-JUL-2003; 2003WO-EP007864
                                                                                                                                                                                                                                             26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Obeta coat protein mutant which can used as part of the repetitive
                                                                             Bachmann MF,
                                                                                                                                                                                                                                                                             WO2004016282-A1
                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteriophage Obeta.
                                                                                                             (CYTO-) CYTOS BIOTECHNOLOGY AG (NOVS ) NOVARTIS PHARMA AG.
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                                                                                Tissot
                                                                                                                                                                                                                                                                                                        /note= "Wild type Lys
                                                                                                                                                                                                                                                                                                                                     /note= "Wild type Lys substituted by Arg"
                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                                                Ortmann
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Composition comprising a core particle with at least one attachment site,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes a novel composition comprising a virus-like core particle with at least one attachment site, and an antigenic amyloid beta CC 1-6 peptide. The new composition comprises: a core particle with at least cone first attachment site; and at least one antigen or antigenic cone first attachment site; and at least one antigen or antigenic cone antigenic determinant is a Amyloid beta 1-6 peptide, and where the second attachment site comprises: an attachment site not naturally cocurring with the antigen or antigenic determinant; or an attachment site is the naturally occurring with the antigen or antigenic determinant. The second attachment site is capable of association to the first attachment site and the beta 1-6 peptide and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful for the manufacture of a medicament for treating Alzheimer's disease and related diseases. This is the amino acid sequence of an RNA bacteriophage Obeta coat protein mutant that can be used in the proparation of the compositions and vaccines of the invention.
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Matches 132; Conserv
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04-MAY-2001; 2001US-0288549P.
05-OCT-2001; 2001US-032698P.
07-NOV-2001; 2001US-0331045P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory;
cytostatic; antiviral; antidiabetic; hypoglycaemic; antigen array;
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10-DEC-2002
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h as Alzheimer's disease
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Pred. No. 8.2e-71;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to a novel ordered and repetitive antigen array CC used in the production of vaccines for infectious diseases. The invention CC also discloses a composition comprising a non-natural molecular scaffold CC comprising a core particle selected from a core particle of a non-natural CC origin and a core particle of natural origin and an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond. Also disclosed is an CC antigen or antigenic determinant with at least one second attachment site, where the second attachment site, where the antigen or antigenic determinant is amyloid beta peptide CC (Abetal-42) or its fragment and where the second attachment site is selected from an attachment site not naturally occurring with the antigen or antigenic determinant, where the second attachment site is capable of association through at least one non-peptide bond to the first attachment site and where the antigen or antigenic determinant and the capacitation to form an ordered and comparise antiquent array. The invention also comprises a coat protein capable of forming a capsid which comprises mutant Obeta coat protein capable of forming a capsid which comprises main acid sequences (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C capable of forming a capsid which comprises minate (C c capable of forming the comprise of the invention
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                   Molecular antigen array; vaccine; antigen; antimicrobial; molecular scaffold, amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgE-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus erythematosus; osteoporosis; inflammatory immune disease; mymphadenopathy; hultiple sclerosis; immunoproliferative disease; lymphadenopathy; Alzheimer's disease; mangioimmunoproliferative lymphadenopathy; immunoblastive lymphadenopathy; anguotative lymphadenopathy; immunoblastive lymphadenopathy; humanoblastive lymphadenopathy; humanoblastive lymphadenopathy; humanoblastive lymphadenopathy;
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                                                                                                                                                                                                                                                                                             Bacteriophage Q-beta Al coat protein mutant Qbeta-240
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Core particle of a non-natural origin; (1) a core particle selected from: (1) a core particle of an organiser comprising at least one first attachment conjuin; and (2) a core particle of antural origin; and (2) a core particle of antural conjuin; and (2) a core particle of antural conjuin; and (2) a core particle by at least cone site, where the organiser comprising at least one first attachment connected to the core particle by at least one ocovalent bond; (b) an antigen or antigenic determinant with at least cone second attachment site, where the antigen or antigenic determinant; and where the second attachment site not naturally occurring with the antigen or antigenic determinant; and (ii) an attachment site is capable of association concurring with the antigen or antigenic determinant; and (ii) an attachment site is capable of association concurring with the antigen or antigenic determinant; and (ii) an attachment site; and through at least one non-peptide bond to the first attachment site; and (ii) an attachment site; and (ii) 
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04-MAY-2001; 2001US-0288549P.
05-OCT-2001; 2001US-032698P.
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MAURER P.
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Tissot A,
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                          The present invention describes a composition (C) for enhancing an immune cresponse against an antigen or a virus-like particle in an antigen, or a C comprises a virus-like particle (VLP) bound to at least one antigen, or a C VLP capable of being recognised by the immune system of the animal. Also described: (1) enhancing an immune response sgainst an antigen or a VLP in an animal comprising introducing (C) into the animal; (2) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; (3) immunising or treating an animal comprising administering the vaccine; and (4) enhancing anti-viral protection in an animal comprising introducing (C) into the animal. (C) the animal by administering the vaccine; and (4) enhancing anti-viral protection in an animal comprising introducing (C) into the animal. (C) has cytostatic, virucide, antibacterial, antiparasitic, fungicide, antihlergic, immunosuppressive, antiaddictive, antiinflammatory, antihrevaccines can be considered antiallabetic, neuroprotective, nootropic, osteopathic, antiparasitic antihreumatic and antiarthritic activities. (C) or the vaccines can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory; antithyroid; antidiabetic; neuroprotective; nootropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis; Alzheimer's disease; osteoporosis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A composition, useful for enhancing or a virus-like particle, enhancing or treating tumors or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-SEP-2001; 2001US-0318967P
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   immune response against an antigen or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     an immune response against an antigen anti-viral protection in an animal, infectious diseases, e.g. viral
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                                                                                                                                                                                                                                                                                                                                                                                                                                                05-OCT-2001; 2001US-032699BP.
07-NOV-2001; 2001US-0331045F).
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-IB000166.
19-JUL-2002; 2002US-0396637P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction; congestive heart failure; kidney failure; retinal haemorrhage; mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABU09689 standard;
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Pred. No. 1.9e-70;
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New angiotensin peptide moiety carrier conjugate comprising a an angiotensin peptide moiety, useful for treating or preventi disorder associated with renin-activated angiotensin, e.g. hyperical states of the second sec

2003-430264/40.

Claim 20;

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93-94;

97pp;

invention describes an angiotensin peptide moiety carrier conjugate

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Search completed: January 4, 2005, 09:11:01
Job time: 39.6824 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder angiotensin useful with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kidney failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta coat protein used in the preparation of the vaccine conjugates of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 132 AA;
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                                                                                                                          121 LIDAIDQLNPAY 132
                                                                                                                                                                                                          61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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Perfect score:
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

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6: /cgn2_6/ptodata/1/iaa/backfIles1.pep:*
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PCT-US95-14377-3
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24620, A
44620, A
44620, A
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1182, App
1185, App
216, Appl
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2745, Appl
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RESULT 2 US-09-948- ; Sequence ; Patent N ; GENERAL ; APPLICE; ; APPLICE; ; TITLE C	90 VQ	Query M Best Lo Matches Qy	RESULT 1 US-09-485-7 ; Sequence ; Patent No ; GENERAL I ; APPLICAN ; APPLICAN ; AFFLICAN ; TILE OF ; TILE OF ; FILE REF ; CURRENT ; CURRENT ; PRIOR AP ; PRIOR FI ; PRIOR TO ; NUMBER O ; SOFTWARE ; SEQ ID NO ; LENGTH: ; TYPE: P ; ORGANIS ; FEATURE ; OTHER I US-09-485-7	44444333320 8900144444 8444498
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US/09948722 efan H. E. n nerculosis Vaccine	1 0	3*; Score 81.5; DH 4; Len 0\$; Pred. No. 0.3; 24; Mismatches 44; InNPRGVNPTNGVASLSQAGA INQNNADIQUVNAISSLTYPGALVKANS	(a 4 a (3 US-09-120-915 3 US-09-134-001 3 US-08-781-891 4 US-09-618-639-4 4 US-09-976-594 54 US-09-206-942 4 US-09-206-942 4 US-09-206-942 4 US-09-206-942 5 US-09-206-942 6 US-09-206-942 7 US-09-206-942 8 US-09-206-942 9 US-09-308-32 9 US-08-720-484 9 US-08-730-484 9 US-09-398-239
v	NPTACTANGSCDESVTROKYADVT 	5; DB 4; L 0.3; ches 44; ASLSQAGA :: :	ccine 85,717 14.7 05109 Artificial Sequence:	915-7 -001C-2885 -1961-76 -156-75 -594-41 -411-13 -942-35 -942-35 -942-37 -942-26 -942-37 -942
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	FTQYSTD 102 AKIDYD 252	; Gaps 5; VPALEKRV 48 : LPVKRDSL 197	ant	tce 7, Applice 7, Applice 76, Applice 76, Applice 11, Applice 13, Applice 35, Applice 26, Applice 26, Applice 12, Applice 12, Applice 12, Applice 5, Appli

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US-09-328-352-4249
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Best Local Similarity 25.2
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LENGTH: 409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4249, Application US/09328352
Patent No. 6562958
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: HUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BANGANNII FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: GTC99-03PA
                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 8252
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PRIOR FILING DATE: 1997-08-23
NUMBER OF SEQ ID NOS: 2
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PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: PCT/EP98/05109
PRIOR FILING DATE: 1998-08-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 100564-00079
CURRENT APPLICATION NUMBER: US/09/948,722
CURRENT FILING DATE: 2002-04-08
                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: recombinant nu OTHER INFORMATION: molecule comprising a domain of Mycobacterium and OTHER INFORMATION: escape domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
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                                                               170 IAYLRETGFTNNAALVPENTAKPYSEITF-----EEVMESVKTIAHMLKASKQILDDL
                                                                                                                                      118 RGGTRLNVNVPRNALISFAVNPVDGTTPI----IAKPNORLTIRDLLAPGRTGSN---A
                                  126 DQL 128
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                                                                                                66 VKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDBERAFVRTELAALLASPLLIDAI 125
                                                                                                                                                                         13 RDGKQTLVLNPR------GVNPTNGVASLSQAGAVPALEKRVTV-SVSQPSRNRKNYKVQ 65
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PQL 225
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; Sequence 44620,
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                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 23637
LENGTH: 916
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Best Local S
Matches 27
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GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: MACC J. Rubenfield et al.
APPLICANT: MACC J. RUBENFIELD AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TO DEFERENCE: 107196.136
TO DEFERENCE: 107196.136
                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                      Query Match
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SEQ ID NO 16691
LENGTH: 818
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATITIES OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/074,788
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PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 1998-07-27
                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Pseudomonas aeruginosa
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                                                                                                                            56 SRNRKNYKVQVKIQNPTAC---TANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRT 110
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                                                                                                                                                                                                        9 GNIGRDG-----KQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVS-----QP 55
                                                                                                                                                                                                                                              27; Conservative
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                                                                                          EERGAEYRAQESAHREHPCRSIEANADCPLSADAGKW-----PWTIFSDSATRWALRT 901
                                                                                                                                                                  GLLGGEGALARURAAIVVQARRIDPFQGLAAALPVGGLPALEQAEDFAFGGSLDQRWVQA 848
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  Application US/09270767
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rubenfield et al.
NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
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BER: US 60/074,788
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;; Pred. No. 10;
17; Mismatches
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Pred. No. 8.
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; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-44620
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TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 44620
LENGTH: 502
  TELEPAX: (212) 940-8636
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 AMINO
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GENERAL INFORMATION:
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Patent No. 5534257
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Best Local Similarity 24.2%;
                                                                                                                                                                                                        SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/090,148
FILING DATE: 08/11/93
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9101550.3
FILING DATE: 01/24/91
APPLICATION UNMBER: PCT/GB92/00124
FILING DATE: 01/22/92
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APPLICANT: Mastic
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                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
COMPUTER: IBM PS2-486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                            REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
                                                                                                                                                    NAME: Nissenbaum, Israel REGISTRATION NUMBER: 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 575 Mac
CITY: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         315 AXXVVKYFSCTSSATRHSTRCISSYVSQSLGPSPLLGXADXKDFVRTLXPLATFISSPNL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   258 TIAVN---ISPSPSPTPQPIVEPVEQVENSVTVTVASPEVPPVAXARNDRXRQTRSTIKN 314
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18 TLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQP-----SRNRKNYKVQVKIQN 70
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AMINO
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Stockley, Peter George
Talbot, Simon John
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                                                                                                                                                                                                                              PCT/GB92/00124
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Pred. No. 5.
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US-09-302-626B-182
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; DATE: 1972
US-08-090-148-1
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                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 195
SOFTWARE: PatentIn Ver. 2
SEQ ID NO 182
                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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Patent No. 6709660
                                                                                                                                      Matches
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Best Local
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CURRENT FILING DATE: 1999-04-30
PRIOR APPLICATION NUMBER: PCT/IB99/00103
PRIOR FILING DATE: 1999-01-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Pizza, Mariagrazia
APPLICANT: Grandi, Guido
TITLE OF INVENTION: Meningococcal Antigens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: CHIR0159
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Scalato,
                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                   LENGTH: 300
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PUBLICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AUTHORS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE: gene coding for the TITLE: bacteriophage MS2 coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AUTHORS: Min Jou, W.; Haegeman, G. AUTHORS: Ysebaert, M.; Fiers, W. TITLE: Nucleotide sequence of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: NOT RELEVANT
64 ADKSAPKNQQAVILQTANGLPQVNIQTPS-----SQGVSVNR----
                               50 VSVSQPSRNR-----KNYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDE 103
                                                                                                     10 NIGRDGKQTLVLNPRGVNPTNGVASLSQAGAV-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27 NPTNGVA-----SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVKIQNPTACTANGSC
                                                                    5 NVHRDGKSMQDSEAASVRVT-GAASVSSARAAFGFRMAAFSVMLALGVAAFSPAPASGII 63
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Rappuoli, Rino
                                                                                                                                      Conservative
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                                                                                                                                                       10.4%; Score 69; I
23.2%; Pred. No. 3
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                                                                                                                                          Mismatches
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                                                                                                                                        51; Indels
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 ----FKQFDVDE 108
                                                                                                                                          58;
                                                                                                       -PALEKRVT 49
                                                                                                                                        Gaps
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                                                                                                        NUMBER OF SEQ ID NOS: 1387
SOFTWARE: CuraPatSeqFormatter Version
SEQ ID NO 216
LENTH: 766
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 216, Application Patent No. 6753314
GENERAL INFORMATION:
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 195
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 185
LENGTH: 622
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APPLICANT: C
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CURRENT APPLICATION NUMBER: US/09/302,626B
CURRENT FILING DATE: 1999-04-30
PRIOR APPLICATION NUMBER: PCT/IB99/00103
PRIOR FILING DATE: 1999-01-14
                                                                                                                                                                                              FILE REFERENCE: 15966-542
CURRENT APPLICATION NUMBER: US/09/538,092
CURRENT FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: 60/127,352
PRIOR FILING DATE: 1999-04-01
PRIOR APPLICATION NUMBER: 60/178,965
PRIOR APPLICATION NUMBER: 60/178,965
PRIOR FILING DATE: 2000-02-01
                                                                                                                                                                                                                                                                                                                                             APPLICANT: Giot, Loic
APPLICANT: Mansfield, Traci A.
TITLE OF INVENTION: Protein-Protein Complexes and Method of Using Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Pizza, Mariagrazia
APPLICANT: Grandi, Guido
APPLICANT: Grandi, Guido
APPLICANTION: Meningococcal Antigens
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NAME/KEY: misc_feature LOCATION: (0)...(0)
                                               TYPE: PRT
ORGANISM: Saccharomyces
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23 NVHRDGKSMQDSEAASVRVT-GAASVSSARAAFGFRMAAFSVMLALGVAAFSPAPASGII 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10 NIGROGKQTLVLNPRGVNPTNGVASLSQAGAV------
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Similarity 23.2%;
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Rappuoli, Rino
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                                                                      cerevisiae
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US-09-206-942-49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Haemophilus influenzae US-09-206-942-49
                                                                                                                                                                                                                  US-09-206-942-47
                                    GENERAL INFORMATION:
APPLICANT: Loosmore, Sheena M.
APPLICANT: Yang, Yan-Ping
APPLICANT: Klein, Michel H.
TITLE OF INVENTION: Protective Recombinant Haemophilus Influenzae High
TITLE OF INVENTION: Molecular Weight Proteins
FILE REFERENCE: 1038-861 MIS:jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 49
LENGTH: 1073
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Best Local 9
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Best Local Similarity
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APPLICANT: Yang, Yan-Ping
APPLICANT: Klein, Michel H.
TITLE OF INVENTION: Protective Recombinant Haemophilus Influenzae High
TITLE OF INVENTION: Molecular Weight Proteins
FILE REFERENCE: 1038-861 MIS:jb
CURRENT FILING DATE: 1998-12-08
EARLIER APPLICATION NUMBER: 05/167,568
EARLIER FILING DATE: 1998-10-07
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    CURRENT
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                                                                                                                                                                                                                                                                                          880 ISA 882
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APPLICATION NUMBER: US/09/206,942 FILING DATE: 1998-12-08
                                                                                                                                                                                                                                                                                                                                                                                                           55 PSRN----RKNYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        54 QPSRNRKNYKVQVKIQNPTACTANGSCDPSVT-----RQKYADVTFSFTQYSTDEER 105
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 TVTLGNI---GRDGKQTLVLNPRGVNPTNGVASLSQAGAV-----PALEKRVTVSVS 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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Pred. No.
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28;
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RESULT 14
US-09-045-186-2
Sequence 2, Application US/09045186
; Sequence 1087154
; PATENT NO. 6087154
; GENERAL INFORMATION:
; APPLICANT: Baez, Melvyn
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US-09-198-452A-745
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LENGTH: 381
TYPE: PRT
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Matches
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APPLICANT: Griffais,
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Best Local S
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TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis,
TITLE OF INVENTION: and treatment of infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EARLIER FILING DATE: 1998-10-07
NUMBER OF SEQ ID NOS: 95
SOFTWARE: PatentIn Ver: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/198,452A CURRENT FILING DATE: 1998-11-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Chlamydia pneumoniae
FEATURE:
RAME/KEY: SITE
LOCATION: 1...381
OTHER INFORMATION: Xaa=unknown or other
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27;
                                                                                                                                                                                                                                                                                  63 KVQVKIQNPTACTANGSCDPSVTRQKYADVTFS-----FTQYSTDEERAFVRTELAALL 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 55 PSRN----RKNYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTE 111
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                                                                                                                                                                                                                                                KHNLRDEKLAACPKNSAASLSAKKSHTKKTTPGSIPSKVFSKFDATQDKTFQKTSGSAFP 116
                                                                                                                                                                         AKPTTLKELEE 127
                                                                                                                                                                                                                                                                                                                       VKTVIL-NIGRKILQGIKXXKKKIGIXSGLFFLDLV-----LLGVSXQRPTETSANV
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6849
                                                                                                                                                                                                                                                                                                                                                                                                               10.3%; Score 68.5; DI
19.8%; Pred. No. 6.2;
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                                                                                                                                                                                                                                                                                                                                                                                              27; Mismatches
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Pred. No.
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28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 381;
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; MOLECULE TYPE: protein US-09-045-186-2
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                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 5571695
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/232,144
FILING DATE: 26-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 276-0756
TELEFAX: (317) 276-3861
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 383 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Human Neuropeptide NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: GAYLO, PAUL J.
REGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: P-:
                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                        APPLICANT: SELBIE, Lisa
APPLICANT: HERZOG, Herbert
APPLICANT: SHINE, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cates,
                                                                                                                                                                                                                                 STREET: 555 lstn
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: CLASSIFICATION:
                                                                                                                                                                             COUNTRY: US
ZIP: 20004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CLASSIFICATION:
                                                                                                                                                                                                                                                                               ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16 KQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK
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Lill Corporate Center
                                                                                                                                                                                                                                                        E: Rothwell, Figg,
555 13th St, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            United States of America
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: RHESUS NEUROPEPTIDE Y1 RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10.3%; Score 68.5; DI 25.0%; Pred. No. 6.3;
                                     US/08/232,144
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Suite 701-East
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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       seq length: 0
seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-622-064-8
664
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            110:
111:
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119:
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/cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

1 2 3 3 4 4 6 6 7 7 7 11 11 12 13	Result No.
00000000000000000000000000000000000000	Score
100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 99.5	Query
132 132 132 132 132 132 132 132 132 132	Length
11111111111111111111111111111111111111	Bd
US-10-243-739-25 US-10-244-065-25 US-10-289-454-25 US-10-050-902-257 US-10-050-898-257 US-10-346-190-25 US-10-346-190-25 US-10-486-25 US-10-289-456-25 US-10-622-064-8 US-10-622-064-8 US-10-622-064-8 US-10-622-064-9 US-10-622-087-19 US-10-622-087-19 US-10-243-739-23 US-10-244-065-23	ID
Sequence 25, Appl Sequence 25, Appl Sequence 25, Appl Sequence 27, App Sequence 27, App Sequence 25, Appl Sequence 16, Appl Sequence 25, Appl Sequence 19, Appl Sequence 19, Appl Sequence 23, Appl Sequence 23, Appl Sequence 23, Appl	Description

45	44	43	42	41	40	39	38	37	36	35	34	ω ω	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	1
652	655	655	655	655	655	655	655	655	655	655	655	658	658	658	658	658	658	658	658	658	658	658	661	661	661	661	661	661	661	661	
98.2	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	99.1	99.1	99.1	99.1	99.1	99.1					99.1		•	•		٠	•		•	
132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	132	
14	16	15	15	15	15	14	14	14	14	14	14	16	15	15	15	15	14	14	14	14	14	14	16	15	15	15	15	14	14	14	
-10-243-739-	6	-10-622-124-	-10-622-064-	-10-289-456-	-10-465-811-	6-190-2	-10-050-898-2	-10-050-902-	-10-289-454-	-10-244-065-	3-739-	-10-622-087-	-10-622-124-	-10-622-064-	US-10-289-456-27	-811-	-10-346-190-	-10-050-898-25	-10-050-902-2	-10-289-454-2	-10-244-065-	243-	-10-622-087-	-10-622-124-	622-064-	-289-	5-8	0-346-190-	-25	-10-	
O	Sequence 20,	e 20,	9,	e 26,	e 17,	e 26,	e 259,	e 259,	e 26,	e 26,	e 26,	e 21,	21,	10,	e 27,	e 18,	e 27,	258	e 258	27,	27,	27,	17	17	6	23	e 14	23	e 25	e 25	י
Appl	Appl	Appl	Appli	Appl	Appl	Appl	_	_	App1	Appl	Appl	App1	Appl	Appl	Appl	Appl	Appl	, App		Appl	Appl	App1	Appl	, Appl	Appli	Appl	Appl	Appl	, App	, App	,

ALIGNMENTS

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US-10-243-739-25
; Sequence 25, Application US/10243739
; Sequence 25, Application No. US20030091593A1
; GENERAL INFORMATION:
   APPLICANT: Bachmann, Martin F.
   APPLICANT: Storni, Tazio
   APPLICANT: Lechner, Franziska
   APPLICANT: Lechner, Franziska
   TITLE OF INVENTION: Inmune Responses Induced by Virus Like Particles
   FILE REFERENCE: 1700.0210001
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; TYPE: PRT
; ORGANISM: Bacteriophage Q-beta
US-10-243-739-25
                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-9-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 25
                                                                                                                                                                                                  Matches 132;
                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
121 LIDAIDQLNPAY 132
                                                                     61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                         61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
                                                                                                                     Conservative
                                                                                                                                                                                                  100.0%; Score 664; DB 14; 100.0%; Pred. No. 2.1e-66; ... Mismatches 0;
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                                                                                                                                                                                                                                          Length 132;
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APPLICANT: Schwarz, Katrin
APPLICANT: Meijerink, Edwin
APPLICANT: Lipowsky, Gerad
APPLICANT: Pumpens, Paul
APPLICANT: Pumpens, Paul
APPLICANT: Renhofa, Regina
ITITE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Particl
ITITE OF INVENTION: Method of Preparation and Use
FILE REFERENCE: 1700.0220001
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 25
LENGTH: 132
TYPE: PRI
COGANISM: Bacteriophage Q-beta
US-10-244-065-25
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Matches
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APPLICANT: Scorni, Ta
APPLICANT: Maurer, Pe
APPLICANT: Tissot, Al
APPLICANT: Schwarz, I
APPLICANT: Meijerink,
                                                                                                                                                                                                                                                                                                                Sequence 25, Application US/10289454 Publication No. US20030157479A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                     APPLICANT: Bachmann, Martin
APPLICANT: Jennings, Gary
APPLICANT: Jennings, Gary
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
                                                                                                                      CURRENT APPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
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les 132; Conserv
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Maurer, Patrick
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; NUMBER OF SEQ ID NOS: 386

SOFTWARE: PatentIn version 3.2

; SEQ ID NO 25

; LENGTH: 132

; TYPE: PRT

; ORGANIEM: Bacteriophage Q-beta-250

US-10-289-454-25
                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Qb
US-10-050-902-257
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SEQ ID NO 257
LENGTH: 132
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Best Local Similarity
Matches 132; Conserv
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Best Local Similarity
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CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-05-04
PRIOR PILING DATE: 2001-10-05
PRIOR FILING DATE: 2001-10-05
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TITLE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0190004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 350
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                                      121 LIDAIDQLNPAY 132
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                                                                                                             61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDBERAFVRTELAALLASPL 120
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LIDAIDQLNPAY 132
                                                                                  NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
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Sebbel, Peter
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100.0%; Pred. No. 2.1e-66;
htive 0; Mismatches 0;
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100.0%; Pred. No. 2.1e-66;
tive 0; Mismatches 0;
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RESULT 5 US-10-050-898-257

Sequence 257, Appublication No.

Application US/10050898 o. US20030175711A1

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Sequence 25, Application US/10346190
Publication No. US20030219459A1
GENERAL INFORMATION:
APPLICANT: Bachmann, Martin
APPLICANT: Maurer, Patrick
APPLICANT: Maurer, Patrick
APPLICANT: Renner, Wolfgang A.
TITLE OF INVENTION: Prion Protein Carrier-Conjugates
FILE REFERENCE: 1700.029003
CURRENT APPLICATION NUMBER: US/10/346,190
CURRENT APPLICATION NUMBER: 60/396,590
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/389,898
PRIOR APPLICATION NUMBER: 60/389,898
PRIOR APPLICATION NUMBER: 60/389,898
PRIOR APPLICATION NUMBER: 60/389,898
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TITLE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0190005
CURRENT APPLICATION NUMBER: US/10/050,898
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 60/289,549
PRIOR APPLICATION NUMBER: US 60/26,998
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
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Matches
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NUMBER OF SEQ ID NOS: 350
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 257
LENGTH: 132
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-10-050-898-257
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APPLICANT:
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Tissot, Alain
Maurer, Patrick
Lechner, Franziska
Sebbel, Peter
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Ortmann, Rainer
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; LENGTH: 132
; TYPE: PRT
; ORGANISM: Qb :
US-10-346-190-25
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; OTHER INFORMATION: Bacteriophage Q-beta 250 mutant
US-10-465-811-16
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PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR FILING DATE: 2002-01-18
NUMBER OF SEQ ID NOS: 164
SOFTWARE: PatentIn version 3.1
SEQ ID NO 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16, Applica Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.2
SEQ ID NO 16
LENGTH: 132
TYPE: PRT
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                                                                                                                                                                                                                                                  Matches 132;
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Best Local Similarity
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APPLICANT: RENNER, WOLFGANG A
TITLE OF INVENTION: PACKAGED VIRUS-LIKE PARTICLES FOR USE AS ADJUVANTS:
TITLE OF INVENTION: METHOD OF PREPARATION AND USE
PILE REFERENCE: 1700.0290004
CURRENT APPLICATION NUMBER: US/10/465,811
CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: US 60/389,898
PRIOR PILING DATE: 2002-06-20
PRIOR PILING DATE: 2002-06-20
NUMBER OF SEQ ID NOS: 131
                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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                                       121 LIDAIDQLNPAY 132
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No. US20040005338A1
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                                                                                                                                                                                                                                                                                           DB 15;
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RESULT 8 US-10-289-456-25 ; Sequence 25, Application US/10289456

GENERAL INFORMATION

APPLICANT: Bachmann,
APPLICANT: Maurer, I
APPLICANT: Spohn, G

Maurer, Patrick Spohn, Gunther

Martin

Publication No. US20040033211A1

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CURRENT APPLICATION UNMBER: US/10/289,456
CURRENT FILING DATE: 2002-11-07
PRIOR APPLICATION UNMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR FILING DATE: 2002-01-21
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 170
SOFTWARE: PATENTIN UNS: 170
SOFTWARE: PATENTIN UNS: 170
SOFTWARE: PATENTIN UNS: 170
COMMANDER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 170
SOFTWARE: PATENTIN UNS: 170
COMMANDER: US 60/331,045
PRIOR TILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 170
SOFTWARE: PATENTIN UNS: 170
COMMANDER: US 60/331,045
PRIOR TILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 170
SOFTWARE: PATENTIN UNS: 170
COMMANDER: US 60/331,045
PRIOR TILING DATE: 2001-11-07
NUMBER: US 60/331,045
PRIOR TILING DATE: 2001-11-07
NUMBER: US 60/331,045
PRIOR FILING DATE: 2002-07-19
PRIOR FILING DATE: 2002-0
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                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Bacteriophage Q-beta 250 mutant
US-10-622-064-8
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APPLICANT: Bachmann, Martin F
APPLICANT: Maurer, Patrik F
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Publication No. US20040059094A1
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Best Local S
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,575
PRIOR FILING DATE: 2002-07-18
NUMBER OF SEQ ID NOS: 33
SOFTWARE: Patentin version 3.2
SOFTWARE: Patentin version 3.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Maurer, Patrik F
TITLE OF INVENTION: Hapten-Carrier Conjugates and Uses Thereof
FILE REFERENCE: 1700.0300001
                                                                                                                                                                                                                                                                                                                                        LENGTH: 132
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Local Similarity 100.0%; (
hes 132; Conservative 0;
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132; Conserv
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1 ARLETVILGNIGRDGKQTLVLNPRGVNPINGVASLSQAGAVPALEKRVIVSVSQPSRNRK 60
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                                                                                                                   100.0%; Score 664; DB 15; larity 100.0%; Pred. No. 2.1e-66; Conservative 0; Mismatches 0;
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Pred. No. 2.1e-66;
; Mismatches 0;
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PRIOR FILING DATE: 2002-07-19
NUMBER OF SEQ ID NOS: 146
SOFTWARE: PatentIn version 3.2
SEQ ID NO 19
LENGTH: 132
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                          APPLICANT: Staufenbiel, Matthias
APPLICANT: Frey, Peter
TITLE OF INVENTION: Amyloid Beta 1-6 Antigen Arrays
FILE REFERENCE: 1700.0350002
CURRENT APPLICATION NUMBER: US/10/622,087
CURRENT FILING DATE: 2003-07-18
FRIOR APPLICATION NUMBER: US 60/396,639
PRIOR PILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: US 60/470,432
PRIOR APPLICATION NUMBER: US 60/470,432
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NUMBER OF SEQ ID NOS: 93
SOFTWARE: PatentIn version 3.2
SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19, Application US/10622087 Publication No. US20040141984A1
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Best Local Similarity
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TITLE OF INVENTION: Ghrelin-Carrier Conjugates
FILE REFERENCE: 1700.0340001
CURRENT APPLICATION NUMBER: US/10/622,124
CURRENT FILING DATE: 2003-07-18
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Publication No. US20040076645A1
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APPLICANT: Tissot, Alain
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                                                                                                                                                                                                                                                                                                      APPLICANT: Ortmann, Rainer
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PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
LENGTH: 132
TYPE: PRT
ORGANISM: Bacteriophage Q-beta
US-10-243-739-23
                                                                                                                                 RESULT 13
US-10-244-065-23
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Publication No. US20030091593A1

GENERAL INFORMATION:

APPLICANT: Bachmann, Martin F.

APPLICANT: Storni, Taxio

APPLICANT: Lechner, Franziska

TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of
TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
                                                                          Sequence 23, Applica Publication No. US20 GENERAL INFORMATION:
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Best Local Similarity
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Best Local
APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Maurer, Patrick
APPLICANT: Tissot, Alain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 1700.0210001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
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                                                                                          Application US/10244065
to. US20030099668A1
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Pred. No. 4.5e-66;
1; Mismatches 0
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Pred. No. 2.1e-66;
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APPLICANT: Cielens, Indulis
APPLICANT: Renhofa, Regina
ITITLE OF INVENTION: Packaging of Immunostimulatory Sul
ITITLE OF INVENTION: Method of Preparation and Use
FILE REFERENCE: 1700.0220001
CURRENT APPLICATION NUMBER: US/10/244,065
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR FILING DATE: 2002-04-22
PRIOR FILING DATE: 2002-04-22
PRIOR FILING DATE: 2001-09-14
PRIOR FILING DATE: 2001-09-14
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; ORGANISM: Bacteriophage Q-beta
US-10-244-065-23
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                                                                                      US-10-289-454-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23, Applica Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
LENGTH: 132
Query Match
Best Local Similarity
Matches 131; Conserv
                                                                                                                                                                   SOFTWARE: PatentIn version 3.2 SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local
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PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 386
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bachmann, Martin
APPLICANT: Jennings, Gary
APPLICANT: Sonderegger, Ivo
APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
CURRENT APPLICATION NUMBER: US/10/289,454
                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PT/IB02/00166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 73
                                                                                                                              TYPE: PRT
                                                                                                       ORGANISM: Bacteriophage Q-beta-240
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No. US20030157479A1
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Lipowsky, Gerad
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                      99.5%;
Score 661; DB 14;
Pred. No. 4.5e-66;
1; Mismatches 0;
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Pred. No. 4.5e-66;
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                                          Length 132;
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Gaps
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1 ARLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60

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APPLICANT: Renner, WOLLYGRAY A.
APPLICANT: Bachmann, Martin
APPLICANT: Hachmann, Martin
APPLICANT: Hechmer, Patrick
APPLICANT: Lechmer, Pranziska
APPLICANT: Lechmer, Pranziska
APPLICANT: Piossek, Christine
ITILE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0190004
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-01-9
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR APPLICATION NUMBER: US 60/31,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 350
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 255
LENGTH: 132
TYPE: PRT
ORGANISM: QB 240
US-10-050-902-255
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US-10-050-902-255
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Search completed: January 4, 2005, 09:41:22 Job time: 32.7351 secs
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Best Local Similarity
Matches 131; Conserv
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APPLICANT: Renner, Wolfgang A.
APPLICANT: Bachmann, Martin
APPLICANT: Tissot, Alain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 255, Appublication No.
                                                                                     121
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b. US20030175290A1
                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                           99.5%; Score 661; DB 14; Length 132; 99.2%; Pred. No. 4.5e-66; tive 1; Mismatches 0; Indels
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Title:
Perfect score:
Sequence:
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Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                         Total number of hits satisfying chosen parameters:
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                                                                                                                                          BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                      January 4, 2005, 09:02:51; Search time 7.88938 Seconds (without alignments) 1609.836 Million cell updates/sec
                                                                                                 283416 seqs, 96216763 residues
                                                                                                                                                                                                       1 ARLETVTLGNIGRDGKQTLV.....AALLASPLLIDAIDQLNPAY 132
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664
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Copyright (c) 1993 - 2005 Compugen Ltd.
                                                               283416
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. pir_79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Database :

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SUMMARIES

Result No.	Score	Query Match	Length	DΒ	ID	Description
_	652	98.2	132	-	VCBPQB	coat protein - pha
N	526.5	79.3	331	N	S01964	through pro
ω	116	17.5	131	_	VCBPP1	dg - r
4.	87.5	13.2	530	N	S22340	igeriolysir
u	85.5	12.9	540	N	T00646	hypothetical prote
0	81.5	12.3	529	N	S24231	м
7	81.5	12.3	529	N	A43505	listeriolysin O pr
80	81.5	12.3	529	N	AC1100	0
9	81	12.2	830	ผ	857537	ب
10	80.5	12.1	1502	H	RGBYH1	377
11	77.5	11.7	528	N	S22341	ivanolysin precurs
12	77	11.6	432	N	T31660	hypothetical prote
13	76.5	11.5	136	N	C98221	
14	76.5	11.5	136	N	AE3065	
15	76.5	11.5	1052	N	C64221	hypothetical 114K
16	76.5	11.5	1861	N	T13845	381
17	76	11.4	130	,_	A46324	coat protein - pha
18	76	11.4	282	N	AI0186	7
19	75.5	11.4	1097	N	AD2572	hypothetical prote
20	75	•	130	<u>ب</u>	VCBPGA	
21	74.5	11.2	130	_	VCBPFR	ı
22	74.5		161	4.	I55480	na
23	74	11.1	340	N	S18650	
24	74	11.1	340	N	A42008	homeotic protein H
25	73.5	11.1	520	_	ACMSD1	nicotinic acetylch
26	73.5	11.1	601	N	A55921	serine/threonine k
27	73.5		719	N	T39271	conserved hypothet
28	73	11.0	243	Н	VHVUPT	nucleocapsid prote
29	73	11.0	430	N	G88884	protein K09B11.10

RESULT 2 801964 201964 readthrough protein - phage SP C;Species: phage SP C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004

ALIGNMENTS

B &	g Q	D 64	Que Bes Mat	A, Residu A, Residu A, Residu J. Biol. A, Title: A, Title: A, Access A, Access A, Molecu A, Residu C, Superf	A;Molecu A;Residu A;Cross- R;Stoll, J. Biol. A;Title: A;Refere A;Access	RESULT VCBPQB COAT p C;Spec C;Date C;Acce R;Esca R;Bio A;Titl A;Refe A;Acce
121 LIDAIDQLNPAY 132 121 LIDAIDQLNPAY 132	61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120 	1 ARLETVILGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60 -	Query Match 98.2%; Score 652; DB 1; Length 132; Best Local Similarity 97.7%; Pred. No. 5.2e-57; Matches 129; Conservative 2; Mismatches 1; Indels 0; Gaps 0;	A;Residues: 1-60 (STO)-A;Residues: 1-60 (STO)-A;Residues: 1-60 (STO)-A;Residues: 1-60 (STO)-A;R:Maita, T.; Konigsberg, W. R;Maita, T.; Konigsberg, W. Biol. Chem. 246, 5003-5024, 1971 A;Ritle: The amino acid sequence of the Qbeta coat protein. A;Reference number: A92088; MUID:71288580; PMID:5570434 A;Recession: A92088 A;Recession: L920, Drotein A;Residues: 1-21, Drotein A;Residues: 1-21, Drotein A;Residues: 1-21, Drotein C;Superfamily: phage GA coat protein	A;Molecule type: mRNA A;Residues: 1-80 <esc> A;Residues: 1-80 KESC> A;Cross-references: UNIPROT: P03615 R;Stoll, B:; Wilson, K.J.; Reiser, J.; Weissmann, C. T. Biol. Chem. 252, 990-993, 1977 J. Biol. Chem. 252, 990-993, 1977 A;Title: Revised amino acid sequence of Qbeta coat protein between positions 1 and 60. A;Reference number: A92221; MUID: 77118576; PMID: 838709</esc>	RESULT 1 VCBEQB VCBEQB Coat protein - phage Q-beta C;Species: phage Q-beta C;Species: phage Q-beta C;Date: 29-Uul-1981 #sequence_revision 24-Sep-1981 #text_change 09-Uul-2004 C;Accession: A92240; A92221; A92088; A04224 R;Bacarmis, C.; Sastry, P.A.; Billeter, M.A. J. Biol. Chem. 253, 8390-8399, 1978 J;Title: Determination of the first half of the coat protein cistron of bacteriophage Qbe A;Reference number: A92240; MUID:79048469; PMID:361741 A;Accession: A92240

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seeligeriolysin - Listeria seeligeri
C;Species: Listeria seeligeri
C;Date: 22-Nov-1993 #sequence_revision 01-Dec-1995
C;Accession: S22340
R;Haas A; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete sequence of A;Reference number: S22340; MUID:92182018: PMTD.154
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Dhaese, P.; Vandekerckhove, J.S.; Van Montagu, M.C.
Eur. J. Biochem. 94, 375-386, 1979
A;Title: The primary structure of the coat protein of the
A;Reference number: A04225; MUID:79148387; PMID:107028
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A;Status: preliminary
A;Molecule type: genomic RNA
A;Residues: 1-331 <INO>
A;Cross-references: UNIPROT: P09677; EMBL:X07489
A;Note: the authors translated the readthrough
C;Superfamily: phage GA coat protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 42
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Best Local S
Matches 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oat protein - phage PRR1
;Species: phage PRR1
;Date: 30-Nov-1979 #sequence_revision 30-Nov-1979 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Accession: A04225
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                                                                                                                                                                                                                                                             DAIDQLNPAY 132
                                                                                                                                                                                                                                                                                                                             KIQNPT--ACTANGSCDPSVTRQKYADVTFSFTQYSTDEER-AFVRTELAALLASPLLI-
                                                                                                                                                                                                                                                                                                                                                                                                          QTLVLNPRGVNPT------NGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQV
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                                                                                                                                                                                                                                                                                                 KLVVPVVQSQTVNGIVTPVVVRTSYVTVDFDYDARSTTKERNNFVGMIADALKADLMLVH 121
                                                                                                                                                                                                                                                                                                                                                                     QNLVLKDREATPNDHTFVPRDIRDNVGEVVESTGVPIGESRFTISLRKTSNGR--YKSTL
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77.3%;
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 s: complete sequence of ilo
MUID:92182018; PMID:1543752
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Pred.
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Mismatches 15
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                                                                                            09-Jul-200'
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hypothetical protein F316.9 - Arabidopsis thaliana (;Species: Arabidopsis thaliana (mouse-ear cress) c;Pate: 01-Feb-1999 #sequence_revision 01-Feb-1999 #text_change 09-C;Accession: T00646 R;Federspiel, N.A.; Palm, C.J.; Conway, A.B.; Kurtz, D.B.; Conway, Vysotskaia, V.S.; Yu, G.; Ecker, J.; Theologis, A.; Davis, R.W. submitted to the EMBL Data Library, February 1998 A;Reference number: Z14197 A;Accession: T00646
A;Reference number: S24230;
A;Accession: S24231
A;Status: nucleic acid seque
A;Molecule type: DNA
                                                                                  C;Species: Listeria monocytogenes
A;Variety: strain 12067
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995
C;Accession: S24231
R;Rasmussen, O.F.; Beck, T.; Olsen, J.E.; Dons, L.;
Infect. Immun. 59, 3945-3951, 1991
                                                                                                                                                                                                                  RESULT
S24231
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A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-530 <HAA>
A;Cross-references: UNIPROT:P31830; EMBL:X60462;
A;Note: the authors translated the codon GCC for C;Superfamily: dipeptide transport protein
                                                                                                                                                                                                                                                                                                                                                         В
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C; Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-540 <FED>
A;Cross-references: UNIPROT:O48683; EMBL:AC002396; NID:g2749918; PIDN:AAC00577.1;
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T00646
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                                                                       A; Title: Listeria monocytogenes isolates can be classified into
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A; Introns: 14/3;
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                                                                                                                                                                                              listeriolysin precursor - Listeria monocytogenes (strain 12067)
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Best Local
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Best Local
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30; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                      IGRDGKQTLVLNPRG----VNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKN-YKVQV
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                                                                                                                                                                                                                                                                                       KPAAPMSKSPQGFATPRVYKPAPTKTSLSTSHSSLKKEK----
                                                                                                                                                                                                                                                                                                                      KIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pyruvate
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                    acid sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                             12.9%; Score 85.5;
26.1%; Pred. No. 1.
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                                                      MUID:92040062;
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Pred. No. 1;
                     shown;
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residue 287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 540;
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C;Species: Listeria monocytogenes
C;Date: 21-Oct-1992 #sequence_revision 21-Oct-1992 #text_change 09-Jul-2004
C;Date: 21-Oct-1992 #sequence_revision 21-Oct-1992 #text_change 09-Jul-2004
C;Accession: A43505; S05306; A47606; S12400; A61079
R;Mengaud, J; Vicente, M.F.; Chenevert, J.; Pereira, J.M.; Geoffroy, C.; Gicquel-Sanzey
Infect. Immun. 56, 766-772, 1988
Infect. Immun. 56, 766-772, 1988
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C;Superfamily: dipeptide transport protein
C;Keywords: virulence factor
F;1-25/Domain: signal sequence #status predicted <
F;26-529/Product: listeriolysin #status predicted
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A;Experimental source: strain 12067, serotype 4b
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1991
C;Genetics:
                                                                                                                       A;Residues: 413-480 <ME2>
A;Cross-references: GB.M29171
R;Mtchel, E; Reich, K.A.; Favier, R.; Berche, P.; Cossart, P.
Mol. Microbiol. 4, 2167-2178, 1990
Mol. Microbiol. 4, 2167-2178, 1990
A;Title: Attenuated mutants of the intracellular bacterium Listeria monocytogenes obtain A;Reference number: S12400; MUID:91211627; PMID:1965218
A;Accession: S12400
                                                                                                                                                                                                                                                                                                                                                                           Infect. Immun. 55, 3225-3227, 1987
A,Title: Identification of the structural gene encoding A,Reference number: A47606; MUID:88057627; PMID:2824384
A,Recession: A47606
A,Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: EMBL:X15127, NID:g44106; PIDN:CAA33223.1; PID:g44107 A;Experimental source: strain EGD A;Note: this sequence is derived from a weakly hemolytic strain, serotype R;Mote: this sequence is derived from a weakly hemolytic strain, serotype R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Domann, E.; Chakraborty, T.
Nucleic Acids Res. 17, 6406, 1989
A;Title: Nucleotide sequence of the listeriolysin gene from a Listeria monocytogenes ser A;Reference number: S05306; MUID:89366684; PMID:2505236
A;Accession: S05306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Title: Expression in Escherichia coli and sequence analysis of the listeriolysin O det A;Reference number: A43505; MUID:88153053; PMID:3126142 A;Accession: A43505
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A; Experimental source: C; Genetics:
                                                              A; Molecule type: DNA
A; Residues: 483-493 <MIC>
                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
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A; Residues: 1-529 < DOM>
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A;Note: this sequence is derived from a strongly hemolytic strain, serotype 1/2c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-529 <MEN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      49
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                             strain LO28,
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                             serotype 1/2c
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S57537

MKTI protein - yeast (Saccharomyces cerevisiae)

N.Alternate names: protein N2302; protein YNL085w

C;Species: Saccharomyces cerevisiae

C;Species: Saccharomyces cerevisiae

C;Date: 10-Oct-1995 #sequence revision 10-Nov-1995 #text change 09-Jul-2004

C;Accession: S57537; S50279; S63024; S63017; S65096

C;Accession: S57537; S50279; S63024; S63017; S65096

R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M.

submitted to the EMBL Data Library, June 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT:P13128; GB:NC_003210; PIDN:CAD00729.1; PID:g16409567; GSPDB:(A;Experimental source: strain EGD-e C;Genetics: A;Gene: hly C;Superfamily: dipeptide transport protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; Maitournam, A.; Matok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; Voss, H.; Wehland, A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Accession: AC1100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Listeria monocytogenes
C;Date: 27-Nov-2001 #sequence_rev
C;Accession: AC1100
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C;Keywords: virulence factor
E;1-25/Domain: signal sequence #status predicted <SIG>
F;26-529/Product: listeriolysin O #status predicted <M
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A; Residues: 1-529 < GLA>
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A; Reference number: S57533
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kdgneyivvekkkkksinqnnadiqvvnaissltypgalvkanselvenqpdvlpvkrdsl 152
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22.0%; Pred. No. 4;
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A;Description: The sequence
A;Reference number: S59386
A;Accession: S59400
A;Molecule type: DNA
A;Residues: 1-1502 <JOH>
A;Cross-references: EMBL:U20
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A;Molecule type: DNA
A;Residues: 1-830 <SOL>
A;Cross-references: UNIPROT:P40850; EMBL:X89016; NID:g887621; PID:g887626
A;Cross-references: UNIPROT:P40850; EMBL:X89016; NID:g887621; PID:g887626
                                                                                                                                             R;Johnson, D.

R;Johnson, D.

submitted to the EMBL Data Library,

submitted to the sequence of S. c
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A;Status: nucleic acid s
A;Molecule type: DNA
A;Residues: 1-830 <SGP>
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R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M.

Yeast 12, 485-491, 1996

A;Title: The sequence of a 17 933 bp segment of Saccharomyces cerevisiae chromosome A;Reference number: S65092; MUID:96310628; PMID:8740422
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A;Residues: 1-29, 'G', 31-808, 'TMKTCLNYH' <VER>
A;Residues: 1-29, 'G', 31-808, 'TMKTCLNYH' <VER>
A;Cross-references: EMBL:U09129; NID:g520475; PIDN:AAC49470.1;
R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J:P.G.; Remacha, M.
submitted to the Protein Sequence Database, April 1996
A;Reference number: S63018
A;Accession: S63024
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A;Title: Sequence of MKT1, needed for propagation of M(2) A;Reference number: S50279; MUID:95176705; PMID:7532890 A;Accession: S50279
                                                                                                                                                                                                                                    N;Alternate names: protein L9672.1; protein YLR256w; regulatory protein CYP: C;Species: Saccharomyces cerevisiae C;Decies: Saccharomyces cerevision 23-Feb-1996 #text_change 12-Nov-1999 C;Date: 30-Sep-1991 #sequence revision 23-Feb-1996 #text_change 12-Nov-1999 C;Accession: S59400; A31312; S15447; S05804; S15446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Map position: 14L
C;Superfamily: Saccharomyces cerevisiae MKT1 protein
C;Keywords: transmembrane protein
F;615-631/Domain: transmembrane #status predicted <TMM>
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A; Accession: S63017
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: SGD:S0005029; MIPS:YNL085w
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Best Local S
Matches 26
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;Residues: 1-830 <SOW>
;Cross-references: EMBL:Z71361; NID:g1301982; PID:g1301983; MIPS:YNL085w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Molecule type: DNA
Residues: 569-830 <POE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65 QVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEER 105
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      EMBL: U20865;
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      NID:g662330;
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                                                                                                                                                      cerevisiae
                                                                                                                                                                                                                                                                                                                         protein YLR256w; regulatory protein CYP1; regulator
                                                                                                                                                                                                                                                                                                                                                          yeast (Saccharomyces cerevisiae)
                                                                                                                                                                              February
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      PIDN:AAB67387.1; PID:g662331; GSPDB:GN0001
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A;Map position: 12R
C;Superfamily: regulatory protein HAP1; GAL4 zinc binuclear cluster C;Superfamily: regulatory protein HAP1; GAL4 zinc binuclear cluster C;Superfamily: heme binding; transcription regulation; zin E;1-148/Domain: DNA binding #status predicted <DNA>
F;9-98/Domain: GAL4 zinc binuclear cluster homology <GAL4>
F;64-84/Region: zinc finger CCCC motif
F;177-189/Region: glutamine-rich
F;177-189/Region: glutamine-wich
F;245-445/Domain: heme binding #status predicted <HEM>
F;249-344,323-326,347-352,373-378,389-394,415-420/Region: 6-residue
F;1308-1481/Domain: activation element #status predicted <ACT>
F;1388-1481/Region: acidic
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A;Title: Functional dissection and sequence of yeast HAP1 activator A;Reference number: A31312; MUID:89106221; PMID:2643482
A;Accession: A31312
A;Accession: S36683
A;Molecule type: DNA
A;Residues: 1-319,'T',321-528 <KRE>
A;Cross-references: EMBL:X60461
C;Genetics:
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                                                                                                                                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-528 < HAS>
                                                                                                                                                                                                                                                                                                                           A,Title: Listeriolysin genes: complete sequence of ilo A,Reference number: S22340; MUID:92182018; PMID:1543752 A,Accession: S22341
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R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ivanolysin precursor - Listeria ivanovii
C;Species: Listeria ivanovii
C;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change
C;Accession: S22341; S36683
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A;Molecule type: DNA
A;Rosidues: 1-62,'R',64-1305,'Y',1306-1470,'LVDFYRADFPIWE'
A;Rosidues: EMBL:X13793
A:Cross-references: EMBL:X13793
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R;Pfeifer, K.; Kim, K.S.; Kogan, S.; Guarente,
                                                                                                                                                                                             R;Kreft,
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                                                                                                                                      A; Reference number:
                                                                                                                                                                  submitted to the EMBL Data
                                                                                                                                                                                                                    A; Note: the authors translated the
                                                                                                                                                                                                                                                   A;Cross-references: UNIPROT:P31831; EMBL:X60461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references:
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20.6%; Pred. No. 17;
                                                                                                                                                               Library, July 1991
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zinc finger
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C;Specie
C;Date:
C;Access
R;Bird,
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C;Superfamily: dipeptide transport protein
F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-528/Product: ivanolysin #status predicted <MAT>
                                                                                                                                                                                                                             R;Goodner, B.; Hinkle, G.; G
A.; Liu, F.; Wollam, C.; Al
Science 294, 2323-2328, 2001
                                                                                                                                                                                                                                                                                                      hypothetical protein AGR_L_1428 [imported] - Agrobacterium tumefaciens (strain C;Species: Agrobacterium tumefaciens C;Species: Agrobacterium tumefaciens C;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Accession: T31660
R;Bird, A.P.; Clark, V.; Jones, S.J.; Leitgeb, S.; submitted to the EMBL Data Library, December 1996
A;Reference number: Z21049
A;Accession: T31660
                                                                                                A; Cross-references:
                                                                                                                  A; Molecule type: DNA
A; Residues: 1-136 < KUR>
                                                                                                                                                                       A;Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrobacterium tun
A;Reference number: A97359; MUID:21608551; PMID:11743194
A;Accession: C98221
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein COS41.6 - sea squirt (Ciona intestinalis)
C;Species: Ciona intestinalis
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 09-Jul-2004
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                                                                                                                                                     Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DTSSEGEVTKDGPGNLAENP---TPSNAARELQESVASEVLETTVVKSAIQ-EQDSSAYR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RDGKQTLVL-----NPRGVNPTNGVASLSQAGA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                 KE-NPONAPACPNRNGNCVSTNSNKTPNDSKIEIVDNSDDNTTDEDTQIDTNE 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TVSVSQPSRNRKNYKVQVKIQNPTACTANGSCDPSVTR-----QKYADVTFSFTQYSTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                              UNIPROT:Q8U8F3; GB:AE007870;
                                                                                                                                                                                                                                                  G.; Gattung, S.; Miller, N.; Blanchard, M.; Qurollo, C.; Allinger, M.; Doughty, D.; Scott, C.; Lappas, C.;
                                    chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11.6%;
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21.2%;
11.5%;
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Pred. No. 8.7;
17; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 77.5;
Pred. No. 9.
  Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL: Z83760; NID: e1014349; PID: e289996; PIDN: CAB0609
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  76.5;
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  DB
                                                                                              PIDN: AAK89293.1;
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  ν,
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  Length
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                                                                                              PID:g15159127;
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                                                                                                                                                                                                                                                B.; Goldm
Markelz,
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                                                                                                                                                                                                                                                    Goldman,
kelz, B.;
                                                                                                GSPDB:C
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C;Species: Mycoplasma genitalium
C;Date: 17-Nov-1995 #sequence revision 17-Nov-1995 #text_change 09-Jul-2004
C;Accession: C64221; JQ0092; S18702; S18703
R;Fraser, C.M.; Gocayne, J.D.; White, O.; Adams, M.D.; Clayton, R.A.; Fleis M.; Fuhrmann, J.; Nguyen, D.; Utterback, T.R.; Saudek, D.M.; Phillips, C.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8.
                                                                                               A;Status: nucleic acid sequence not shown; translation not A;Molecule type: DNA
A;Rosidues: 1-1052 <TIGR>
A;Cross-references: UNIPROT: P22747; GB:U39696; GB:L43967;
A;Experimental source: strain G-37
A;Experimental source: Strain G-37
A;Innaine, J.M.; Loechel, S.; Collier, A.M.; Barile, M.F.;
                                                                                                                                                                                                                                                        C.A.; Venter, J.C. Science 270, 397-403, 1995 Science 270, 397-403, 1995 A;Title: The minimal gene complement of Mycoplasma genitalium. A;Reference number: A64200, MUID:96026346; PMID:7569993 A;Accession: C64221
A;Title: Nucleotide sequence of the MgPa (1A;Reference number: JQ0090; MUID:90060815; A;Accession: JQ0092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 15
C64221
hypothetical 114K protein (MgPa 3' region) - Mycoplasma genitalium
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C;Genetics:
A;Gene: Atu4139
A;Map position: linear chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-136 <KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Authors: Yoo, H.; Tao, ster, E.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; erage, G.; Gillet, W.; Grant, C.; Guenthmer, D.; F.; Karp, P.; Romero, P.; Zhang, S. Science 294, 2317-2323, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            conserved hypothetical protein Atu4139 [imported] - C;Species: Agrobacterium tumefaciens C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 # C;Accession: AE3065
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Best Local (
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                                                                            259-267, 1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11 IGRDGKQTLVLNPRGVNPTNGVASLSQAGAV-----PALEKRVTVSVSQPSRNR
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24.6%; Pre
---ive 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24.6%; Pred. No. 2.5; tive 16; Mismatches
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                                                                                                                                                                                                                                     not shown; translation not shown
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Pred. No. 2.5;
6; Mismatches
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                              (mgp) operon of 
5; PMID:2583522
                                                                                                                                                                                                                                                                                                                                                                                              Adams, M.D.; Clayton, R.A.; Fleischmann,
T.R.; Saudek, D.M.; Phillips, C.A.; Merri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, L. D.; Kutyavin, T.; Levy, R.; Li, M.; McClella
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PIDN:AAL44939.1; PID:g17742593; GSPDB:G1
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                                                                                                       M.F.; Hu,
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A;Residues: 1-1052 <INA>
A;Cross-references: GB.M31431; NID:g150157; PIDN:AAA25421.1; PID:g150160
A;Experimental source: strain G-37 (ATCC 33530)
R;Peterson, S.N.; Schramm, N.; Hu, P.; Bott, K.F.; Hutchison III, C.A.
Nucleic Acids Res. 19, 6027-6031, 1991
A;Title: A random sequencing approach for placing markers on the physical map of Mycopla A;Reference number: S18693; MUID:92051396; PMID:1945886
A;Accession: S18702
A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA
A;Residues: 902-964 <PET>
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, September 1991
C;Genetics:
A;Genetic code: SGC3
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Search completed: January 4, 2005, 09:17:30 Job time: 8.03224 secs
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                                                                                                        857 KTVDEIINTPEILNSFFRFTPAF 879
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Result
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Maximum Match 100%
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Maximum DB
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Perfect score:
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Q8VDC2
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064303
COAT BPSP
VAL BPSP
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AAA16663
Q9TOR9
O64307
  Q9L5B9
Q6E942
Q6E9A2
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O64307 enterobacte
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Coat i
                                                      Bacher J.M., Bull J.J., Ellington A.D.; "Evolution of phage with chemically amb: BMC Evol. Biol. 3:24-24(2003).
EMBL; AY099114; AAM33126.1; -.
                                                                                                                      Viruses, ssRNA positive-strand viruses, no Allolevivirus, Allolevivirus subgroup III. NCBI_TaxID=12009;
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EMBL; V00643; CAA23992.1; -.
PIR; A92240; VCBPQB.
PDB; 1QBE; X-ray; A/B/C=1-13:
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                                                                                                                                                              Coat protein.
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                                      SEQUENCE
                                                                                              PubMed=14667253;
                                                                                                       SEQUENCE FROM N.A.
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133 AA;
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IPRO02703; Levi coat.

)1819; Levi_coat; 1.

'"re; Coat protein; Direct protein sequencing; RNA-binding.
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Gene 137:133-137(1993).

EMBL; M909114; AAM31127.1; -.

EMBL; M99039; AAA16663.1; -.

EMBL; M99039; AFA16663.1; -.

EMBL; M99039; F. structural molecule activity; IE;

GO; GO:0019028; C:viral capsid; IEA.
GO; GO:001908; F. structural molecule activity; IE;

InterPro; IPR002703; Levi coat.

InterPro; IPR001904; RNA rec mot.

Pfam; PR01819; Levi coat; 1.

PROSITE; PS00030; RRM_RNP_1; UNKNOWN 1.

SEQUENCE 329 AA; 36134 MW; E0F30215A8C197CB CR
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01-OCT-2002
01-OCT-2002
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Bacteriophage Q-beta.
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Kozlovska T.M., Cielens I., Dreilinna D.,
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PubMed=14667253;
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Q9TOR9;
01-MAY-2000
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                                                                                                          SEQUENCE FROM N.A.

Beekwilder M.J., Nieuwenhuizen R., van Duin J.;

Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.

EMBL; AF059242; AAC14699.1; -.

HSSP; P03615; 1QBE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=94109687; PubMed=7506687; Kozloveka T.M., Cielens I., Dreilinna Ose V., Pumpens P.; Recombinant RNA phage Q beta capsid passembled in Escherichia coli."; Gene 137:133-137(1993).
EMBL; M99039; AAA16663.1; -
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MEDLINE=96190948; PubMed=8609616;
Beekwilder J., Nieuwenhuizen R., Pc
Beekwilder J. Structure model for the
"Secondary structure model for the
Control of A-protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
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Viruses; ssRNA positive-strand
Allolevivirus.
                                  GG; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi coat.
Pfam; PF01819; Levi_coat; 1.
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Nieuwenhuizen R.,
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A1-protein.
Enterobacteria phage M
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                                                                                                                                                          GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi coat.
InterPro; IPR002504; RNA_rec_mot.
Pfam; PF01819; Levi coat; 1.
PROSITE; PS00030; RRM_RNP 1; UNKNOWN 1
SEQUENCE 329 AA; 35880 MW; 372B1DD
                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=95239761; PubMed=7723040; Beckwilder M.J. Nieuwenhuizen R., "Secondary structure model for the RNA phage O beta.";
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases
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MW; 372B1DD5DDB52F15 CRC64;
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                                                                                                                             Score 561;
Pred. No. 2.
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Pred. No. 1e-46;
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Best Local S
Matches 104
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01-AUG-1998 (TrEMBI
01-AUG-1998 (TREMBI
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Alprotein.
Bacteriophage M11.
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GO; G
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"Secondary structure model for the
Control of A-protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
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Beekwilder M.J., Nieuwenhuizen
Beekwilder m.d., nieuwenhuizen
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                                                                                                                    Viruses; ssRNA positive-strand Allolevivirus.
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SEQUENCE 133
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GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi coat.
Pfam; PF01919; Levi_coat; 1.
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     SEQUENCE FROM N.A MEDLINE=95239761;
                                                                                NCBI_TaxID=74336;
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4; Mismatches
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"Secondary structure model for the
Control of A-protein synthesis.";
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GO; GO:0005198; F:structural molecule activity;
InterPro; IPR002703; Levi coat.
InterPro; IPR000504; RNA_rec_mot.
Pfam; PF01819; Levi coat; 1.
PROSITE; PS00030; RRM_RNP_1; UNKNOWN 1.
SEQUENCE 329 AA; 35893 MW; 3E33CD821EE625F4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beekwilder J., Nieuwenhuizen R., Poot R., van
Submitted (MAR-1998) to the EMBL/GenBank/DDBJ
EMBL; AF052431; AAC06251.1; -.
HSSP; P036.15; 10BB.
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RNA phage Q beta.";
J. Mol. Biol. 247:903-917(1995).
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                                                                                    use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                               This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restructe by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                               Hirashima A., Hirose T., Inayama S., "Analysis of the complete nucleotide coliphage SP.";
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                                                               entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                      Nucleic Acids Res. 16:6205-6221(1988).
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                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Viruses; ssRNA positive-strand
Allolevivirus.
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SEQUENCE
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Hirashima A., Hirose T., Inayama S., Inokuchi Y., Jacobson "Analysis of the complete nucleotide sequence of the group coliphage SP.";
Nucleic Acids Res. 16:6205-6221(1988).
-!- FUNCTION: Not yet known.
-!- MISCELLANEOUS: The readthrough protein Al includes the
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01-MAR-1989 (Rel. 10, Last sequence update)
29-MAR-2004 (Rel. 43, Last annotation updat
Readthrough protein A1 [Contains: Coat prot
Bacteriophage SP.
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P09677;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Viruses; ssRNA positive-strand viruses,
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PIR; S01964; S01964.
HSSP; P03615; 1QBE.
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Pfam; PF01819; Levi_coat; 1.
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NYKVQVKIQNPTACTANGSCDPSVTRQKYADVIFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AKLNOVTLSKIGKNGDOTLTLTPRGVNPTNGVASLSEAGAVPALEKRVTVSVAOPSRNRK
                                      AKLNOVTLSKIGKNGDOTLTLTPRGVNPTNGVASLSEAGAVPALEKRVTVSVAQPSRNRK
                                                     ARLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
                                                                                                                                           331 AA;
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                                                                                         Conservative
                                                                                                                                           36203 MW;
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77.3%;
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                                                                                      ; Score 526.5; DB 1;
; Pred. No. 6.8e-43;
14; Mismatches 15;
                                                                                                                                        Coat protein.
Readthrough protein A1.
; 4A6642E4B52C6582 CRC64;
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Pred. No. 2.3e-43;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein].
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IV RNA
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RESULT 12
064310
ID 06431
AC 06431
DT 01-AU
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COS Enter
OC Virus
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Best Local S
Matches 98
                           064310;
01-AUG-1998
01-AUG-1998
01-MAR-2004
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Q9TOR8;
01-MAY-2000
                                                                                                                                                                                                                                                                                     GO; GO:0019028; C:viral capsid; IEA. GO; GO:005198; F:structural molecule InterPro; IPR002703; Levi_coat. Pfam; PF01819; Levi_coat; 1. Coat protein.
                                                                                                                                                                                                                                                                                                                              Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases. BMBL; AR059243; AAC14703.1; -. HSSP; P03615; 1QBE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Beekwilder M.J., Nieuwenhuizen R., "Secondary structure model for the RNA phage Q beta.";
                                                                                                                                                                                                                                                                                                                                                                                                          MEDIINE=96190948; PubMed=8609616; Beekwilder J., Nieuwenhuizen R., Poese structure model for the "Secondary structure model for the Control of A-protein synthesis.";
Viruses; ssRNA
         Al-protein.
Enterobacteria
                                                                    064310
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                                                                                                                    IVDAIDNLNPAY
                            (TrEMBLrel. 07, (TrEMBLrel. 07, (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                               132 AA;
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                                                                    PRELIMINARY;
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         phage
positive-strand
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74.2%; Pred. No. 3.8e-40;
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          NL95.
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13,
24,
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Last
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Last annotation update)
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viruses,
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                            sequence update) annotation update)
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ne first three domains
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last two
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                                                                                                                                                                                                                                                                                                                      activity;
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 DNA stage;
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Best Local
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21-JUL-1986
05-JUL-2004
                                                                                                                                                                                                                                     COAT BPPRR
P03616;
bacteriophage PRR1.";
Eur. J. Biochem. 94:375-386(1979).

-1- FUNCTION: Forms the phage shell; binds to the phag
PIR; A04225; VCBPP1.
InterPro; IPR002703; Levi coat.
Pfam; PF01819; Levi_coat; I.
Coat protein; Direct protein sequencing; RNA-binding.
SEQUENCE 131 AA; 14535 MW; E7E639E1E50FC612 CRC64;
                                                                                                                                                                                                                                                             BPPRR
                                                                                       The primary structure of the
                                                                                                                                                          Viruses; ssRNA positive-strand viruses, unclassified Leviviridae.
                                                                                                                                                                              Coat protein.
Bacteriophage PRR1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi coat.
InterPro; IPR000504; RNA rec_mot.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beekwilder M.J., Nieuwenhuizen R., van Duin J.;
Submitted (APR-1998) to the EMBL/GenBank/DDBJ d.
EMBL; AF059243; AAC14704.1; -.
HSSP; P03615; 1QBE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Beekwilder J., Nieuwenhuizen R., Po
"Secondary structure model for the
Control of A-protein synthesis.";
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                                                                                                               MEDLINE=79148387; PubMed=107028;
                                                                                                                           SEQUENCE
                                                                                                                                               NCBI_TaxID=12024;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Allolevivirus.
NCBI_TaxID=75725;
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fol. Biol. 247:903-917(1995).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PF01819; Levi_coat; 1.
TE; PS00030; RRM_RNP_1; UNKNOWN
NCE 330 AA; 36175 MW; 961E55
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(Rel. 44, Last ann
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Last annotation updat
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Pred. No. 1.1e-39;
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last two domains
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                                                                                                                                                                    stage; Leviviridae;
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                                                       phage RNA
                                                                                         broad-host-range
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RESULT 15
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DT 05-JUL
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DT 05-JUL
DT 05-Mame=1
OS Lister
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Matches 42
                                                               p31830;
01-JUL-1993 (Rel. 26, Creat
01-JUL-1993 (Rel. 26, Last
05-JUL-2004 (Rel. 44, Last
Seeligeriolysin precursor (
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Q8VDC2;
01-MAR-2002
01-MAR-2002
01-OCT-2003
                                               Name=lso;
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STRALN=BALB/c; TISSUE=Liver;
MEDLINE=22350095; PubMed=12461651;
Kiss H., Darai E., Kiss C., Kost-A
  Listeria seeligeri.
Bacteria; Firmicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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Mamm. Genome 13:646-655(2002).
EMBL; AJ428064; CAD20986.1; -.
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Comparative human/murine sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                     47 RVTVSVSQPSRNRKNYKVQV------KIQNPT-----ACTANGSCDPSVTRQ-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TVTLGNIGRDGKQTLV-----LNPRGVNPTNG---VASLSQAGAVP------ALEK
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ysin precursor (Thiol-activated
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Rodentia;
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    Bacillales;
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Pred. No. 0.88
18; Mismatches
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Sciurognathi;
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Matches 33; Conserv
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Pfam; PF01289; Thiol cytolysin; 1.
PRINTS; PR01400; TACTTOLYSIN.
PR0500m; PD007062; Thiol cytolysin; 1.
PR051TE; PS00481; THIOL_CYTOLYSINS; 1.
CYtolysis; Hemolysis; Lipid-binding; Signal; Toxin.
SIGNAL 1 25
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SEQUENCE
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Haas A., Dumbsky M., Kreft J.;
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26 530 Seeligeriolysin.
485 485 Binding to Cholesterol (By similarity).
530 AA; 59181 MW; 416F7A4DD2029866 CRC64;
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1: geneseqp1980s:*
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7: geneseqp2003bs:*
8: geneseqp2004s:*
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This invention relates to a novel ordered and repetitive antigen array used in the production of vaccines for infectious diseases. The invention also discloses a composition comprising a non-natural molecular scaffold comprising a core particle selected from a core particle of a non-natural origin and a core particle of natural origin and an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond. Also disclosed is an antigen or antigenic determinant with at least one second attachment

Claim 18; Page 144; 441pp; English.

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RESULT 1
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Piossek C;
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04-MAY-2001; 2001US-0288549P.
05-OCT-2001; 2001US-032698P.
07-NOV-2001; 2001US-0331045P.
                                                                                                                                                                                                                                                                                Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory; cytostatic; antiviral; antidiabetic; hypoglycaemic; antigen array vaccine; infectious disease.
                                                                                                                                                                                                                                                                                                                                  29-AUG-2003
10-DEC-2002
                                                                                        diseases.
                                                                                              Molecular antigen array used
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                                                                                                                                                                                                                                                                  unidentified bacteriophage.
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                                                                                                                                                        (CYTO-) CYTOS BIOTECHNOLOGY AG.
                                                                                                                                        Bachmann M,
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                                     19-JAN-2001;
04-MAY-2001;
05-OCT-2001;
07-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               molecular scaffold; amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgB-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; grave's disease; systemic lupus erythematosus; osteoporosis; grave's disease; systemic lupus erythematosus; osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiviral, antidiabetic, or hypoglycaemic activities and may be used in immunisation and as a vaccine. The present sequence represents a protein sequence used to create the compositions of the invention. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                         21-JAN-2002; 2002WO-IB000168.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteriophage Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammatory immune disease; myasthenia gravis; multiple sclerosis; immunoproliferative disease lymphadenopathy; Alzheimer's disease; angioimmunoproliferative lymphadenopathy; immunoblastive lymphadenopathy; rheumatoid arthritis; diabetes; infectious disease; factor Xa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecular antigen array; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage Q-beta Al coat protein mutant Qbeta-243.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABG80629 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AKLETVTLGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LIDAIDQLNPAY 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AKLETVTLGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
                                 2001US-0262379P.
2001US-0288549P.
2001US-0326998P.
2001US-0331045P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Qbeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cysteine-containing linker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
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Pred. No. 2.7e-69;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 132;
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ID ABR5

ABR56453 standard; protein; 132

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LIDAIDQLNPAY LIDAIDQLNPAY 132

132

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NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL

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CC molecular scaffold comprising: (i) a core particle selected from: (i) a core particle of a non-natural origin; and (2) a core particle of natural corigin; and (2) a core particle by at least cone second attachment site, is connected to the core particle by at least cone second attachment site, an antigen or antigenic determinant with at least cone second attachment site is selected from: (i) an attachment site not naturally occurring with the antigen or antigenic determinant; and (ii) an extraorment site is selected from: (i) an attachment site not naturally occurring with the antigen or antigenic determinant; and (ii) an extraorment site is capable of association concurring with the antigen or antigenic determinant and the scaffold interact through at least one non-peptide bond to the first attachment site; and through the association to form an ordered and repetitive antigen array. The composition is used in contributed is a process for producing a non-naturally occurring contributed and as a vaccine for diseases such as influenza, graft corrected and repetitive antigen array. The composition is used in court lymphoblastic leukaemia, non-Hodgkin's lymphoma, Grave's disease, systemic lugus erythematosus, inflammatory immune diseases, myasthenia corrected and repetitive antigens, and the scaffold sathma, crave's disease, allergic asthma, constemic such as influenza, graft cost diseases, immunoproliferative lymphadenopathy, immunopathy, anaphylaxis, adult constemic such as influenza, graft cost disease, includence and infectious diseases. The antigens are modified to posses a cleavage site (enterokinase or factor Xa) and a Cysteiner containing N-correction is bacterial protein or peptide which is coupled to the present sequence is bacterial protein or peptide which is coupled to the modified a
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                                                                           Matches 132;
                                                                                                              Query Match
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Renner
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecular antigen array used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-636514/68
                                                                                                                                                   Sequence
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                                                                                               Similarity
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STAUFENBIEL M.
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                                        AKLETVTLGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
   AKLETVTLGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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Bachmann
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                                                                         Conservative
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                                                                                             100.0%;
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M, Tissot A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AG
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                                                                                               Score 663; DB 5;
Pred. No. 2.7e-69;
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                                                                               Mismatches
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Sebbel
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Piossek C;
                                                                                                                Length 132;
                                                                               Indels
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ABR56453;

VLP;

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CC response against an antigen or a virus-like particle in an animal. (C)
CC comprises a virus-like particle (VLP) bound to at least one antigen, or a
CC VLP capable of being recognised by the immune system of the animal. Also
CC described: (1) enhancing an immune response against an antigen or a VLP
CC in an animal comprising introducing (C) into the animal; (2) vaccines
CC comprising (C) together with a pharmaceutical diluent, carrier or
CC excipient; (3) immunising or treating an animal comprising administering
CC the vaccine to the animal, or priming or boosting a T cell response in
CC the animal by administering the vaccine; and (4) enhancing anti-viral
CC protection in an animal comprising introducing (C) into the animal. (C)
CC antiallergic, immunosuppressive, antiaddictive, antiaflammatory,
CC antityroid, antidabetic, neuroprotective, noctropic, osteopathic,
CC antityroid, antidabetic, neuroprotective, noctropic, osteopa
                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory; antithyroid; antiidabetic; neuroprotective; nootropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis; Alzheimer's disease; osteoporosis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or a virus-like purchas, or immunizing or treating tumors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A composition, useful for enhancing an immune response against an antigen or a virus-like particle, enhancing anti-viral protection in an animal, or immunizing or treating tumors or infectious diseases, e.g. viral
                                                                                                                                                                         Alzheimer's disease, osteoporosis, rheumatoid arthritis, or inflau autoimmune disease. ACC69838 to ACC69852 and ABR56401 to ABR56509
                                                                                                     Sequence 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a composition (C) for enhancing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 180-181; 243pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CYTO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage Qbeta.
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                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                presenting cell; APC; immune response; virus like
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                                                                                                                                                  in the exemplification of the present
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Score 663; DB 6;
Pred. No. 2.7e-69;
Mismatches 0;
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AKLETVTLGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60

Matches

132;

0

Indels

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Gaps

0

Best Local Similarity

100.0%;

Score Pred.

663; DB 6; No. 2.7e-69;

Length 132

Query Match

Sequence

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RESULT 4
ABU09690
ID ABU09690
XX ABU0
AX ABU0
AX ABU0
XX Bact
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                                                               The invention describes an angiotensin peptide moiety carrier conjugate comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kidney failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta coat protein used in the preparation of the vaccine conjugates of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-OCT-2001;
07-NOV-2001;
18-JAN-2002;
21-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New angiotensin peptide moiety carrier conjugate comprising a carrier and an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 20; Page 94; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-430264/40
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2001US-0331045P.
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2002WO-IB000166.
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The present invention describes a composition (C) for enhancing an immu response in an animal. (C) comprises a virus-like particle (VLP), and a immunostimulatory substance. The immunostimulatory substance is bound to the VLP, Also described: (1) enhancing an immune response in an animal introducing (C) into the animal; (2) producing (C) for enhancing an immune response in an animal; (3) vaccines comprising (C) together with pharmaceutical diluent, carrier or excipient; and (4) immunising or
                                                                                                                                                                                                                                                                                                                                                                                                      New compositions comprising immunostimulatory substances packaged virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, turns of the compositions of the composition 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-SEP-2002; 2002WO-IB004132.
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22-APR-2002; 2002US-0374145P.
                                                                                                                                                                                                                                                                                                          Disclosure;
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MEIJERINK E.

LIPOWSKY G.

PUMPENS P.

CIELENS I.

RENHOFA R.
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TISSOT A.
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Cielens I,
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                                                                                                                                                                                                                                                                                                   Page 260;
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Renhofa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC treating an animal by: (a) administering the vaccine to the animal; (b) CC priming a T cell response in the animal by administering the vaccine; or CC (c) boosting a T cell response in the animal by administering the vaccine. (C) has immunostimulant, cytostatic, antiallergic, virucide and CC antibacterial activities. (I) can be used in vaccines for enhancing an CC immune response in an animal, particularly a mammal or human. CC immune response in an animal, particularly a mammal or human. CC response, or a cytotoxic T-lymphocyte (CTL) response. (C) or a vaccine CC comprising (C) can also be used for immunising or treating an animal, creptiles or fish. (C) is particularly useful in prophylactic or creptiles or fish. (C) is particularly useful in prophylactic or comprisite traces against allergies, tumours (e.g. breast cancers, CC measles or chicken pox), or bacterial infections (e.g. tuberculosis, CC measles or chicken pox), or bacterial infections (e.g. tuberculosis, catched to the present sequences used in the exemplification of the present invention or syphilis). ACC69790 to ACC69815 and ARR44522 to ABR44612
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                                                                                                                                                                                                                                                                                                                                                          vaccine composition; virus-like particle; core particle; first attachment site; antigen; antigenic determinant; prion prP; PrP peptide; vaccine; neuroprotective; antiinflammatory; prion disease; Bovine Spongiform Encephalopathy; BSE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                       Bacteriophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD24131 standard; protein; 132
                                                                                                                          18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-IB000166.
08-JUL-2002; 2002US-0393725P.
18-JUL-2002; 2002US-0396590P.
                                                                                                                                                                                                                                                                                                                                           Creutzfeldt-Jakob Disease; coat protein; mutant; mutein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JAN-2004
                                                                                                                                                                                                      17-JAN-2003; 2003WO-EP000460.
                                                                                                                                                                                                                                    24-JUL-2003
                                WPI; 2003-598483/56.
                                                              Bachmann M,
                                                                                                                                                                                                                                                                 WO2003059386-A2
                                                                                                                                                                                                                                                                                                Bacteriophage Qbeta
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Similarity 100.0%;
32; Conservative (
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                                                                                               CYTOS BIOTECHNOLOGY AG
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                                                                  Maurer
                                                                                                                                                                                                                                                                                                                                                                                                                                       Qbeta coat protein
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                                                                  Pellicioli
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          mutant Qbeta-243
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No. 2.7e-69;
                                                                   'n
                                                                     WA;
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                                                                                                                                                                                                                                                                                                                                                                                                prion protein;
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A vaccine composition

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Best Local S
Matches 132
                                                                                                                                          07-NOV-2001;
18-JAN-2002;
21-JAN-2002;
19-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to a novel vaccine composition comprising a virus-
like or a core particle with at least one first attachment site and at
least one antigen or antigenic determinant that is a prion protein (PrP)
or its dimer, or a PrP peptide, the antigen or antigenic determinant
being bound to the virus-like or core particle. The vaccine of the
invention may have neuroprotective or antiinflammatory activity. The
composition is useful as a medicament or in manufacturing a medicament
for the treatment or prevention of prion diseases. The prion diseases may
include Bovine Spongiform Encephalopathy (BSE) or Creutzfeldt-Jakob
Disease. The present sequence is the amino acid sequence of a mutant coat
protein from bacteriophage Obeta which may be used during the creation of
the vaccine composition of the invention.
New compositions comprising a core particle and at least one antigen or antigenic determinant, useful for as a vaccine for therapy or prophylax: of bone diseases, particularly mammalian encephalopathies.
                                                                                                                                                                                                                07-NOV-2002; 2002WO-EP012449.
                                                                                                                                                                                                                                            15-MAY-2003.
                                                                                                                                                                                                                                                                        WO2003039225-A2
                                                                                                                                                                                                                                                                                                  Unidentified
                                                                                                                                                                                                                                                                                                                                          osteopathic;
                                                                                                                                                                                                                                                                                                                                                                    Protein for RANKL antigen array to treat bone disease
                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADJ82057 standard; protein; 132 AA
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phage) and at least one prion protein or peptide bound to the virus-like
                                                                                                                 (CYTO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
wes 132; Conser
                                                          2003-441430/41
                                                                                                                                                                                                                                                                                                                              disease;
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                                                                                                                                          ; 2001US-0331045P.
; 2002US-00050902.
; 2002WO-IB000166.
; 2002US-0396635P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                           vaccine;
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                                                                                    Maurer
                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                          vaccine; core particle; antigenic determinant; encephalopathy; immune system stimulation.
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                                                                                                                                                                                                                                                                                                                                                                                                 entry)
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Pred. No. 2.7e-69;
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                prophylaxis
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Best Local
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                                                                                                                                           07-NOV-2001;
18-JAN-2002;
21-JAN-2002;
19-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a composition comprising a core particle having at least one first attachment site, and at least one antigen or antigenic determinant having at least one second attachment site. The antigen or antigenic determinant is a RANKL protein, RANKL fragment or RANKL peptide. The second attachment site is (non-) naturally occurring with the antigen or antigenic determinant, and is capable of association to the first attachment site. The antigen or antigenic determinant and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful as a medicament, or for the manufacture of a medicament for treating bone diseases. The composition is especially useful for as a vaccine for therapy or prophylaxis of bone diseases, particularly mammalian encephalogathies,
                                                                                                                                                                                                                                                                                                                              antiallergic; antiasthmatic; cytostatic; vaccine; virus-like particle; interleukin; II-5; II-13; eotaxin; repetitive antigen array; allergic eosinophilic disease; asthma; Hodgkin's lymphoma.
                                                       WPI; 2003-441518/41.
                                                                                                                                                                                                                  07-NOV-2002; 2002WO-EP012455
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for stimulating mammalian immune system. This sequence represents a
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                                                                                                                                                                                                                                                                                                                                                                                           particle repetitive antigen array peptide #24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the invention.
                                                                                                                                           ; 2001US-0331045P.
; 2002US-00050902.
; 2002WO-IB000166.
; 2002US-0396636P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                 Jennings
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                                                                                                               BIOTECHNOLOGY AG
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                                                                                 Sonderegger
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Pred. No. 2.7e-69;
); Mismatches 0;
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Composition comprising an ordered and repetitive antigen or antigenic determinant array, useful as a medicament, or for manufacturing a medicament for treating allergic eosinophilic diseases, e.g. asthma, or

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SEQ ID NO 24; 245pp; English

Disclosure; SEQ ID NO 15; 252pp; English.

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or chronic viral diseases.

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RESULT 9
ADJ36314
ID ADJ3
XX ADJ363
AC ADJ3
XX ADJ36
AC ADJ3
XX ADJ3
XX ADJ3
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AC ADJ3
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New compositions comprising a virus-like particle (VLP), immunostimulatory substance bound to the VLP, and an antithe VLP, useful for enhancing immune response or for tree
                                                                                                                                                                                                                                                                             Bachman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004000351-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antiallergic; cytostatic; virucide; immunostimulant; vaccine;
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                                                                                                                                                                                      WPI; 2004-108361/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUN-2002; 2002US-0389898P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUN-2003; 2003WO-EP006541.
                                                                                                                                                                                                                                                                                                                                                               (CYTO-) CYTOS BIOTECHNOLOGY AG
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             for treating e.g. tumo:
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The invention relates to a composition comprising a view and at least one antigen, which is a protein or peptide of interleukin CC (II)-5, IL-13 or ectaxin and is bound to the virus-like particle, or a core particle with at least one first attachment site and at least one core partisen with at least one second attachment site, where the antigen is a protein or peptide of IL-5, IL-13 or ectaxin. The second attachment site attachment site and the core is an attachment site naturally or not naturally occurring with the capable of association to the first attachment site, and where the antigen or antigenic determinant and the core particle interact through the association to the first attachment site, and where the antigen or compositions are useful as medicaments, or for manufacturing a medicament cor a vaccine for treating allergic eosinophilic diseases, e.g. asthma, or CC dediction to the compound of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention relates to a composition comprising a virus-like particle at least one antigen, which is a protein or peptide of interleukin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTBLAALLASPL
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ADJ67160
ID ADJ67
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 132;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              an animal comprising a virus-like particle, an immunostimulatory substance bound to the virus-like particle, and an antigen mixed with the virus-like particle. The composition or the vaccine is useful in the manufacture of a pharmaceutical for the treatment of a disorder or disease such as allergies, tumours, chronic diseases and chronic viral diseases. The composition is also useful for enhancing an immune response in an animal. This is the amino acid sequence of a bacteriophage Obeta coat protein mutant, a virus like particle (VLP) that can be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes a composition for enhancing an immune response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 132 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    adjuvant of the invention.
                                                                                                                 New composition comprising a core particle having a first attachment site and an antigen or antigenic determinant which is a ghrelin or ghrelin peptide having a second attachment site, useful for treating obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Phage Qbeta coat protein mutant Qbeta 243 for antigen display array.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADJ67160 standard; protein;
The invention relates to a new composition comprising: (i) a core particle with at least one first attachment site; and (ii) at least antigen or antigenic determinant with at least one second attachment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antigenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           anorectic; core particle;
antigenic array.
                                                                                                                                                                                                                                             Bachmann
                                                                                                                                                                                                                                                                                                                        19-JUL-2002; 2002US-0396638P
                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2004009124-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteriophage Qbeta.
                                                                                Disclosure; SEQ ID NO 18; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                 18-JUL-2003; 2003WO-EP007849
                                                                                                                                                                                                      WPI; 2004-132866/13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>AKLETVTLGKIGKOGKOTLVLNÞRGVNÞTNGVASLSOAGAVÞALEKRVTVSVSOÞSRNRK</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                           Fulurija A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antigenic
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Pred. No. 2.7e-69;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ghrelin; P-pilin;
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Matches 132; Query Match

Similarity

100.0%; Score 663; DB 7; 100.0%; Pred. No. 2.7e-69; ative 0; Mismatches 0;

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Sequence 132 AA;

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LIDAIDQLNPAY 132

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Best Local Similarity
Matches 132; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     site, where the antigen or antigenic determinant is ghrelin or a ghrelin peptide, and where the second attachment site being consisting of an attachment site not naturally occurring with the antigen or antigenic determinant and an attachment site naturally occurring with the antigen or antigenic determinant, where the second attachment site is capable of association to the first attachment site, and where the Shrelin or a ghrelin peptide and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful for treating obesity. The repetitive array may form part of a phage or bacterial display array. This peptide corresponds to a Bacteriophage obeta coat protein mutant which can used as part of the repetitive or
                                                                                                                                                                                                                                                   (CYTO-)
Composition comprising a core particle with at least one attachment site, and an antigenic amyloid beta 1-6 peptide, useful for treating diseases such as Alzheimer's disease.
                                                                                                               WPI; 2004-203731/19.
                                                                                                                                                                                               Bachmann
                                                                                                                                                                                                                                                                                                                                   19-JUL-2002;
15-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                18-JUL-2003; 2003WO-EP007864.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacteriophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuroprotective; nootropic; vaccine; amyloid beta 1-6 peptide;
core particle; antigen array; Alzheimer's disease; RNA bacteriophage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacteriophage Qbeta coat protein mutant N10K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK52194;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADK52194 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 132 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             particle;
protein; C
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2003US-0470432P.
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                                                                                                                                                                                            Tissot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Qbeta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Wild type Asn substituted by Lys"
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                                                                                                                                                                                            ۶
                                                                                                                                                                                               Ortmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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Pred. No. 2.7e-69;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                            Lueoend R,
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RESULT 12
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Best Local
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                                                                                                                                                                                                                       19-JAN-2001;
04-MAY-2001;
05-OCT-2001;
07-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory; cytostatic; antiviral; antidiabetic; hypoglycaemic; antigen array
                                                               Renner WA,
Piossek C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-AUG-2003
10-DEC-2002
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WPI; 2002-627351/67.
                                                                                                                                                                                                                                                                                                                                                                                    21-JAN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unidentified bacteriophage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vaccine; infectious disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pQB240 protein.
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                                                                                                                                                            (CYTO-)
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132; Conserv
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                                                                                                                                                            CYTOS
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                                                                                                                                                                                                                       ; 2001US-0262379P.
; 2001US-0288549P.
; 2001US-0326998P.
; 2001US-0331045P.
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ilarity 100.0%;
Conservative (
                                                                                                Bachmann
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(first entry)
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                                                                                                                                                            BIOTECHNOLOGY AG
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                                                                                             Σ,
                                                                                                Tissot
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Pred. No. 2.7e-69;
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                                                                                             Lechner
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                                                                                             Ŧ,
                                                                                                Sebbel P;
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RESULT 13
ABG94320
ID ABG94
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Query Match
Best Local Sim
Matches 130;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 18; Page 144; 441pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Molecular antigen array used in the production of vaccines for infectious
                                                                                                                                                                  Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory; cytostatic; antiviral; antidiabetic; hypoglycaemic; antigen array
                                                                                                                                                                                                                         pQB251 protein.
                                                                                                                                                                                                                                                               10-DEC-2002
                                                                                                                                                                                                                                                                                                                                                          ABG94320 standard; protein; 132 AA.
                                                                                                          unidentified bacteriophage
                                                                                                                                                vaccine; infectious disease.
                                                                                                                                                                                                                                                                              29-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 655; DB 5;
Pred. No. 2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                  hypoglycaemic; antigen array;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 132;
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21-JAN-2002; 2002WO-IB000166

ABG8063;
ID ABG
XX
AC ABG
XX
AC ABG
XX
DT 29.
XX
DE Bag
XX

ABG80632 standard; protein; 132

Bacteriophage Q-beta Al coat 29-NOV-2002 (first entry)

protein mutant Qbeta-259.

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cc also discloses a composition comprising a non-natural molecular scaffold comprising a core particle selected from a core particle of a non-natural molecular scaffold corigin and a core particle of natural origin and an organiser comprising cart least one first attachment site, where the organiser comprising the core particle by at least one covalent bond. Also disclosed is an cartigen or antigenic determinant with at least one second attachment carticles, where the artigen or antigenic determinant is amyloid beta peptide capable, or its fragment and where the second attachment site is capable of association through at least one non-peptide beta peptide capable of association through at least one non-peptide bond to the first capable of association through at least one non-peptide bond to the first capable of artigen array. The invention also comprises a coat protein capable of forming a capsid which comprises mutant Obsta coat protein capable of forming a capsid which comprises mutant Obsta coat protein antivaral, antidiabetic, or hypoglycaemic activities and may be used in capusce used to create the compositions of the invention. (Updated on 29 AUG-2003 to standardise OS field)
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                                                                                                                                                                                                               Query Match
Best Local Similarity
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04-MAY-2001;
05-OCT-2001;
07-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 18; Page 426; 441pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Molecular antigen array used in the production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-627351/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Renner WA,
Piossek C;
                                                                                                                                                                                                                                                                Sequence 132 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CYTO-) CYTOS BIOTECHNOLOGY AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in the production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention relates
                             121
121
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                                                                                                                                                 AKLETYTLGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRYTVSVSQPSRNRK 60
                             LIDAIDQLNPAY 132
                                                            NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                    NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                                              AKLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
LIDAIDQLNPAY 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; 2001US-0262379P.
; 2001US-0288549P.
; 2001US-0326998P.
; 2001US-0331045P.
                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                 98.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to a novel ordered and repetitive antigen array of vaccines for infectious diseases. The invent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tissot
                                                                                                                                                                                                   <u>,</u>
                                                                                                                                                                                                                 Score 655; DB 5;
Pred. No. 2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P
                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Maurer P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lechner F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of vaccines
                                                                                                                                                                                                                                 Length 132;
                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       molecular scaffold
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for infectious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sebbel P;
                                                                                                                                                                                                   o
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                                                                                                                                                                                                     Gaps
                                                                                                                                   60
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CC molecular scaffold comprising: (i) a core particle selected from: (1) a CC core particle of a non-natural origin; and (2) a core particle of natural CC origin; and (ii) an organiser of comprising at least one first attachment CC site, where the organiser is connected to the core particle by at least one covalent bond; (b) an antigen or antigenic determinant with at least cone covalent bond; (b) an antigen or antigenic determinant is concerned to the core particle by at least cone second attachment site, where the antigen or antigenic determinant is camploid bette apptide (Abetta 1-42) or its fragment, and where the second attachment site naturally occurring with the antigen or antigenic determinant; and the second contribution to the first attachment site; and contribution to the first attachment site; and contributed is a process for producing a non-naturally occurring where the satisfen array. Also included is a process for producing a non-naturally occurring conducted and repetitive antigen array. The composition is used in immunisation and as a vaccine for diseases such as influenza, graft constructed lymphoblastic leukaemia, non-Hodgkin's lymphoma, Grave's disease, coute lymphoblastic leukaemia, non-Hodgkin's lymphoma, Grave's disease, coute lymphoblastic leukaemia, non-Hodgkin's lymphoma, Grave's disease, costeoporosis and infectious diseases. The antigens are modified to posses conscious site (enterokinase or factor Xa) and a Cysteine-containing N-corrections where the second containing N-corrections are the attachment point to a containing N-corrections where the selections are the attachment point to a containing N-corrections.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Maurer
Renner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CYTO-)
(NOVS)
(MAUR/)
(LECH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a composition comprising: (a) a non-natural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Molecular
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05-OCT-2001;
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ORTMANN R.

LUECEND R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STAUPENBIEL M. FREY P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MAURER P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antigen array used
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Bachmann |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 146;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; 2001US-0262379P.
; 2001US-0288549P.
; 2001US-0326998P.
; 2001US-0331045P.
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05-OCT-2001;
07-NOV-2001;
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WPI; 2002-636514/68

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Example 18; Page 144; 418pp; English.
                                                              diseases.
                                                                                       Molecular antigen array used in the production of vaccines for infectious
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CC molecular scaffold comprising: (a) a non-natural CC core particle of a non-atural and CC core particle of a non-atural origin; and (2) a core particle of netural origin; and (2) a core particle of natural CC origin; and (3) a core particle of natural CC origin; and (1) a core particle of natural CC origin; and (1) a core particle of natural CC site, where the organiser is connected to the core particle by at least one covalent bond; (b) an antigen or antigenic determinant with at least CC one second attachment site, where the antigen or antigenic determinant is capturally occurring with the antigen or antigenic determinant; and (ii) an attachment site is selected from: (i) an attachment, and where the second contact through at least one non-peptide bond to the first attachment site; and contact through at least one non-peptide bond to the first attachment site; and contact through the association to form an ordered and repetitive antigen array. The composition is used in contact of an antigenic determinant and the scaffold interact versus host disease, Igs-mediated allergic reactions, anaphylaxis, adult respiratory distress syndrome (ARDS), Crohn's diseases, allergic asthmatic constraints of systemic lymphoblastic leukaemia, non-Hodgkin's lymphoma, Grave's disease, contenuatorid arthritis, diabetes, multiple sclerosis, Alzheimer's disease, contenuatorid arthritis, disease, contenuatorid contenuatorid contenuator

Sequence 132 AA;

Ś 밁 ş Query Match Best Local Similarity Matches 130; 61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120 Н AKLETYTLGKIGKDGKQTLVLNPBGVNPTNGVASLSQAGAVPALEKRYTVSVSQPSRNRK AKLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK Conservative 98.8%; 98.5%; 1; Score 655; DB 5; Pred. No. 2.4e-68; Mismatches Length 132; Indels 0; Gaps 60 60

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4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
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US-09-270-767-44620
US-09-270-767-459B-45
US-08-725-459B-45
US-08-725-459B-42
US-08-725-459B-42
US-08-09-0148-1
US-08-235-836C-11
US-08-235-836C-89
US-09-245-186-2
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US-09-246-186-2
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US-09-248-796A-15-65
US-09-270-673-331
US-09-248-796A-16-65
US-09-270-767-43925
US-09-270-767-43925
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	OTHER INFO 3-09-485-717- Query Match Best Local S Matches 30	PRIOR APPL PRIOR FILI NUMBER OF SOFTWARE: SEQ ID NO 2 LENGTH: 6 TYPE: PRT ORGANISM: FEATURE:	TITLE OF INVENTION: Tube FILE REFERENCE: 16862PUS CURRENT APPLICATION NUMB CURRENT FILING DATE: 20 PRIOR APPLICATION NUMBER PRIOR FILING DATE: 1997-	SULT 1 -09-485-717-2 -09-485-717-2 Sequence 2, Applicat Patent No. 6673353 GENERAL INFORMATION: APPLICANT: Kaufmann		44444	33 3 3 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	332 332 34
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ANG ANG UNIV	Artifi e 84.5 . No. ismatc	.09	ne 717 7		ALIGNMENTS	6360-	61613	618-166 248-796 397-411 248-796 252-991 856-253
SLS SLT SCD	he ci				SIL	-398-239-5 -560-876A-5 -560-876A-6 -843-530B-6	US-08-293-728-2 US-09-421-868-2 US-08-956-171E- US-08-781-986A- US-08-720-484A-	-166-7 -796A- -411-1 -796A- -991A- -253-7
-NPRGUNPTNGVASLSQAGA :: : ONVADIGVVNAISSLTYPGA VKIQNPTACTANGSCDPSVT : : VKNATKSNVNNAVNTLVE	1 1 4;				4	หู้ คือ คือ คือ	-2 -2 -524 -524	47117
RGVNPTNGVASLSQAGAV : :: : :: : :: :	Sequence 44; Len 44; In						5249	-166-76 -796A-16789 -411-13 -796A-20249 -796A-16691 -991A-16691 -253-7
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APPLICANT: Kaufmann, Stefan H. E.
APPLICANT: Hess, Jurgen
TITLE OF INVENTION: Tuberculosis Vaccine

US/09948722

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Best Local Similarity
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PRIOR FILING DATE: 1998-08-12
PRIOR APPLICATION NUMBER: EP 97/114,614.7
PRIOR FILING DATE: 1997-08-23
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                        Query Match
Best Local (
                                                                                                                                                                                                                                                                                          SEQ ID NO 23637
LENGTH: 916
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PRIOR APPLICATION NUMBER: US 09/485,717
PRIOR FILING DATE: 2000-02-22
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CURRENT APPLICATION NUMBER: US/09/948,722
CURRENT FILING DATE: 2002-04-08
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                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/094,190 PRIOR FILING DATE: 1998-07-27 NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: recombinant nucleic acid OTHER INFORMATION: molecule comprising a domain of Mycobacterium and a phagolysomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                        l Similarity
27; Conser
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                                                                                                             GKIGKDG-----KQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVS------QP 55
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EERGAEYRAQESAHREHPCRSIEANADCPLSADAGKW-----PWTIFSDSATRWALRT 901
                                  SRNRKNYKVOVKIONPTAC---TANGSCDPSVTROKYADVTFSFTOYSTDEERAFVRT 110
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                                                                           GLLGGEGALARLRAAIVVQARRIDPFQGLAAALPVGGLPALEQAEDFAFGGSLDQRWVQA 848
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1998-02-18
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22.7%; Pred. No. 0.14
Live 23; Mismatches
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; Sequence 44620, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEO ID NOS: 62517
; SOFTWARE: PatentIN Ver. 2.0
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US-09-270-767-44620
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US-09-328-352-4249
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                                                            US-09-077-098A-6
                                                                                RESULT 6
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APPLICANT: GAIY L. Breton et al.
APPLICANT: GAIY L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
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Patent No. 656
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LENGTH: 409
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Matches
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Sequence 6, Application US/09077098A Patent No. 6544519
GENERAL INFORMATION:
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LENGTH: 502
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                                                                                                                                                                                                                  258 TIAVN---ISPSPSPTPQPIVEPVEQVENSVTVTVASPEVPPVAXARNDRXRQTRSTIKN 314
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24.2%; Pred. No. 6
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APPLICANT: TOKUNAGA, Eiji SAKAGUCHI, Masashi MATSUO, Kazuo

Fukusaburo

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Patent No. 6084068
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COUNTRY: US
ZIP: 75201-6507
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: CONAWAY, RONALD C.
APPLICANT: CONAWAY, JOAN W.
TITLE OF INVENTION: ELONGIN A AND C FUNCTIONAL DOMAINS
NUMBER OF SEQUENCES: 79
                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                             STREET: /-

                                                                                                                                                                                                                                                                                                        ADDRESSEE: SIDLEY & AUSTIN
STREET: 717 N. HARWOOD, SUITE 3400
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FILING DATE: 12-SEP-1997
APPLICATION NUMBER: JP 27,148/1996
APPLICATION NUMBER: JP 27,148/1996
FILING DATE: 19-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: KORNBAU, Anne M.
REGISTRATION NUMBER: 25,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1402
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MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 6:
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APPLICATION NUMBER: US/09/077,098A
FILING DATE: 19-May-1998
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 2042 amino acids
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               62 YKVQVKIQNPTACTANGSCDPSVTRQKYAD--VTFSFTQYSTD 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FTVGVDFTD-TITEGDATDDKKLTTSKSVESYVTKLANFSTD 1443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LDDTTLNKINNPADQDLSNLSESGKNAITGLVDV-----VKKTNSPITVEPSTDSNKKKT 1401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent in Release #1.0, Version #1.30
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TELEFAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/08725459E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: BROWDY AND NEIMARK
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WENTION: NOVEL POLYPEPTIDE FROM HAEMOPHILUS
PARAGALLINARUM AND PROCESS FOR PREPARING THE SAME
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27.2%; Pred. No. 54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13; Mismatches
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RESULT 8
US-08-725-459B-44
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                                                                                                                MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/725,459B
FILING DATE: 04-OCT-1996
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 44, Application US/08725459B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: CONAWAY,
APPLICANT: CONAWAY,
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:NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                  COUNTRY: US
ZIP: 75201-6507
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 31,966
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 214-981-3300
                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 79
CORRESPONDENCE ADDRESS:
ADDRESSEE: SIDLEY & AUSTIN
STREET: 717 N. HARWOOD, SUITE 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION: NAME: HANSEN, EUGENIA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                  REGISTRATION NUMBER: 31,966
REFERENCE/DOCKET NUMBER: 11146/07501
                                                                                                                                                                                                                                                                                                                                                CITY: DALLAS
STATE: TX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity hes 31; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: 1..233
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                     NAME: HANSEN, BUGENIA S. REGISTRATION NUMBER: 31,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/0
FILING DATE: 04-OCT-1996
CLASSIFICATION: 530
                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Peptide LOCATION: 1..233
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STRANDEDNESS: not
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6084068
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                  214-981-3400
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                                214-981-3300
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In Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ELONGIN A AND C FUNCTIONAL DOMAINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RONALD C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JOAN W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "amino acids 202-434 of elegans"
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Pred. No. 2.
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SEQUENCE CHARACTERISTICS

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                                  US-08-725-459B-42
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Query Match
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                                                                                                                                                                                                               TELEFAX: 214-981-3400
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cent No.
                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/725,459B
                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CORRESPONDENCE ADDRESS:
ADDRESSEE: SIDLEY & AUSTIN
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                                                                                                                                MOLECULE TYPE:
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STREET: /-
CTTY: DALLAS
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OTHER INFORMATION:
OTHER INFORMATION:
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PPLICANT: CONAWAY, JOAN W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: not
TOPOLOGY: not rel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ITLE OF INVENTION:
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                                                  LOCATION: 1..434
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: not
TOPOLOGY: not
                                                                                                                                                                                                                                                                                                                  NAME: HANSEN, EUGENIA S. REGISTRATION NUMBER: 31,966
                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 04
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
ZIP: 75201-6507
                                                                                                                                                                                                                                                                 TELEPHONE:
                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
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                                                                                                                                                    TOPOLOGY:
                                                                                                NAME/KEY:
                                                                                 LOCATION:
                                                                                                                                                                                                   ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPSRNRKNYKVQVKIQNPTACTANGSCDPSVTRQK 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  717 N. HARWOOD, SUITE 3400
                                                                                                                                                                                                 434 amino acids
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                                                                                                                                                    not relevant
                                                                                                  Protein
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                                                                                                                                   peptide
                                                                                                                                                                    not relevant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "amino acids 94-434 of C. elegans"
   10.6%;
                                              /note= "entire amino acid sequence
of C. elegans"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13; Mismatches
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   Score 70;
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   DB 3;
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   Length 434;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: GB 91
APPLICATION NUMBER: GB 91
FILING DATE: 01/24/91
APPLICATION NUMBER: PCT/GE
FILING DATE: 01/22/92
ATTORNEY/AGENT INFORMATION:
NAME: NISSENDAUM, ISTAEL
REGISTRATION NUMBER: 27,58
                                                                                                                                                                                                                                                                                                                             TELEFAX: (212) 940-640.
INFORMATION FOR SEQ ID NO: .
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Stockley, Peter Geo
APPLICANT: Talbot, Simon John
TITLE OF INVENTION: Antigen-Pr
TITLE OF INVENTION: Fusion MS2
                                                                                                                                                                                                                                                         TOPOLOGY: NOT MOLECULE TYPE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TUMBER OF SEQUENCES:
                                                                                            PUBLICATION INFORMATION:
                                                                                                                                               LOCATION:
                                                                                                                                                                                   NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: U. FILING DATE: 08/11/93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10022-2585
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                            TYPE: AMINO ACIDS
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                                                                                                                                LOCATION:
                                                                                                                                                                                                                                        ORGANISM:
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                                                                                                             LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     372 QLSQARKNVKIEGKAQLRT-ITPRGGGVPSTSRSR 405
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08090148
Min Jou, W.; Haegeman, G.;
Yesbaert, M.; Fiers, W.
Nucleotide sequence of the
gene coding for the
bacteriophage MS2 coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             575 Madison Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mastico, Robert Allan
                                                                                                                                                                                                                                                                                                                                                                                       (212) 940-6404
                                                                                                                                                                                                                                                                                             NOT RELEVANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM PS2-486
                                                                                                                            represents entire MS2 coat protein in the published
                                                                                                                                                                                                      Coat protein from MS2-RNA
                                                                                                                                                                                                                                          VIRUS
                                                                                                           sequence.
                                                                                                                                                                    Location 1 through 129 below
                                                                                                                                                                                   bacteriophage
                                                                                                                                                                                                                                                                                                                                                                                                          (212) 940-8636
                                                                                                                                                                                                                                                                              PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diskette, 3.5", 1.44Mb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antigen-Presenting Capsid with Fusion MS2-Coat Protein
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US-08-235-836C-11
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                                                                                                                                       Query Match
Best Local Similarity
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APPLICANT: Dunn, John J.
APPLICANT: Luft, Benjami
                                                                                                                                                                                                                                            TELEFAX: (516) 282-372
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 273 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 11973
ZOMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: BN
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 282-7338
                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
FILING DATE: 01-11-93
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APPLICATION NUMBER: US/0
FILING DATE: 29-APR-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PAGES: 0-
                                                                                                                                                                                                                                                                                                                                               NAME: Bogosian, Margaret C.
REGISTRATION NUMBER: 25,324
                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Brookhaven National Laboratory
    239
                                                             184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27 NPTNGVA-----SLSQAGAVPALEKRYTVSVSQPSRNRKNYKVQVKIQNPTACTANGSC
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                                                                                                                                                                                                                               amino acid
KQYTITVKQYDSAGTNLEGT 258
                              KNYKVQVKIQNPTACTANGS 79
                                                          TVTLSKEIAKSGEVTVAL-----NDTNTTQATKKTGAWDSKTSTLTISVNSKKTTQLVFT 238
                                                                                        TVTLGK-IGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR----
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                                                                                                                                                                                                                                           273 amino acids
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                                                                                                                        Conservative
                                                                                                                                                                                                                 linear
                                                                                                                                                                                                protein
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Nr. No. 6248562el Chimeric Proteins Comprising
Nr. Borrelia Polypeptides and Uses Therefor
                                                                                                                                                                                                                                                                                         282-3729
                                                                                                                     10.4%; Score 69; DB
28.8%; Pred. No. 3.6;
tive 12; Mismatches
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                                                                                                                                   DB 3; Length 273;
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                                                                                                                       Indels
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RESULT 13
US-09-045-186-2
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Patent No. 6248562
GENERAL INFORMATION:
APPLICANT: Dunn, John J.
APPLICANT: Luft, Benjamin
TITLE OF INVENTION: No. 67
TITLE OF INVENTION: Borrel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-235-836C-89
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US-08-235-836C-89
                                                                                                                                                                            Sequence 2, Application US/09045186 Patent No. 6087154 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (516) 282-3729
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 273 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/
FILING DATE: 29-APR-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/
FILING DATE: 01-11-93
ATTORNEY/AGENT INFORMATION:
              CORRESPONDENCE ADDRESS:
ADDRESSEB: Eli Lilly and Company
STREET: Lill Corporate Center
CITY: Indianapolis
                                                                                          APPLICANT: Baez, Melvyn
APPLICANT: Cates, Carolyn A.
TITLE OF INVENTION: RHESUS NEUROPEPTIDE Y1 RECEPTOR
NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Bogosian, Margaret C.
REGISTRACIO NUMBER: 25,324
REFERENCE/DOCKET NUMBER: BN
TELECOMMUNICATION INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE SYSTEM: PC-DOS/MS-DOS
COERCEATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 10.4%;
Local Similarity 28.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                        239 KQYTITVKQYDSAGTNLEGT 258
                                                                                                                                                                                                                                                                                                                                                              60 KNYKVQVKIQNPTACTANGS 79
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Indiana
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NN: No. 6248562el Chimeric Proteins Comprisi
NN: Borrelia Polypeptides and Uses Therefor
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Pred. No. 3.6;
12; Mismatches
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US-08-232-144-4
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NAME: Gaylo, Paul J.
REGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: P-11:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 276-0756
TELEPHAX: (317) 276-3861
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
ATTORNEY/AGENT INFORMATION:
NAME: ERNST, Batbara G
REGISTRATION NUMBER: 30,377
REFERENCE/DOCKET NUMBER: 1871
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-783-6040
INFORMATION FOR SEQ ID NO: 4:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: SELBIE, Lisa
APPLICANT: HERZOG, Herbert
APPLICANT: SHINE, John
TITLE OF INVENTION: Human Neuropeptide Y-Y1 Receptor
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 26-MAY-199
CLASSIFICATION: 435
                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
ZIP: 462
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                     STREET: 555
CTTY: Washington
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                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.24
                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/08232144
                                                                                                                                                                                                                                                                                                                                                                    E: Rothwell, Figg, Ernst & Kurz
555 13th St, N.W., Suite 701-East
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              United States of America
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                                                                                                                                                                      US/08/232,144
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; MOLECULE TYPE: protein US-08-232-144-4
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US-08-555-268A-15
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Best Local Similarity
Matches 27; Conserv
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                                                          Matches
                                                             Query Match
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APPLICANT: Bard,
                                                                                                                                                                                                      TELEFAX: (212) 391-052
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
                                                                                                                                STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 44
REFERENCE/DOCKET NUMBER: 44
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP:
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                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY:
                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
16 KOTLVLNPRGVNPTN-----GVA----SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New York
: New York
:RY: U.S.A.
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                                                                                                                                                                                         384 amino acids
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Weinshank, Richard L.
Weinshank, Richard L.
VENTION: DAN ENCODING A HUMAN NEUROPEPTIDE
VENTION: Y/PEPTIDE YY/PANCREATIC POLYPEPTII
VENTION: RECEPTOR (Y4) AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Walker,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                   (212) 278-0400
(212) 391-050
                                                           Conservative
                                                                                                                                               linear
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                                                                                                                                                                                                                                                                                                                                                                       US/08/555,268A
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                                                                                                                                                                                                                          15:
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                                                                                                                                                                                                                                                                                   44743-Z/JPW/MAT
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Pred. No. 7
                                                                         Score 68.5;
Pred. No. 7;
                                                             Mismatches
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Search Job tin	B 8
Search completed: January 4, 2005, 09:35:44 Job time : 11.3622 secs	68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115

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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          seq length: 0 seq length: 2000000000
Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*

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Gapop 10.0 , Gapext 0.5
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

13	12	11	10	9	8	7	δ	ຫ	4	ω	N	_	Result No.
655	655	663	663	663	663	663	663	663	663	663	663	663	Score
98.8	98.8	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	Query Match L
132	132	132	132	132	132	132	132	132	132	132	132	132	Length DB
14	14	16	15	15	15	15	14	14	14	14	14	14	H
US-10-243-739-26	US-10-243-739-23	US-10-622-087-18	US-10-622-124-18	US-10-622-064-7	US-10-289-456-24	US-10-465-811-15	US-10-346-190-24	US-10-050-898-256	US-10-050-902-256	US-10-289-454-24	US-10-244-065-24	US-10-243-739-24	ID
	Sequence 23, Appl	Sequence 18, Appl	Sequence 18, Appl	7		Sequence 15, Appl	Sequence 24, Appl		Sequence 256, App	Sequence 24, Appl	Sequence 24, Appl	Sequence 24, Appl	Description

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14	14	14	14	14	14	14	14	14	14	14	14	16	16	15	15	15	15	15	15	15	15	14	14	14	14	14	14	14	14	14
0-050-902-	-10-050-902-	-10-050-902-15	-10-289-454-	-10-289-454-	-10-289-454-	-10-244-065-	-10-244-065-	-10-244-065-	43-739-	0-243-739-	-10-243-739-	22-087-	0-622-087-	-10-622-124-	-10-622-124-	-10-622-064-	-10-622-064-	-10-289-456-	-10-289-456-2	-10-465-81	-10-465-811-1	-10-346-190-	US-10-346-190-23	-10-050-8	0-050-898-2	-10-050-902-25	-10-050-902-25	-10-289-454-2	-10-289-454-2	-10-244-065-2
258	257	159	27,		10,	27,	e 25,	e 10,	e 27,	e 25,		e 20,	17,	e 20,	e 17		ģ	26	e 23	17,	e 14,	e 26,	e 23,	e 259,	e 255,	e 259	e 255	e 26	e 23,	e 26,

ALIGNMENTS

Q B Q	Query Ma Best Loo Matches Qy	RESULT 1 US-10-24 Sequer Sequer Publia GENERA APPLIA APPLIA TITLE TITLE TITLE TITLE TITLE TITLE TITLE TITLE TITLE SOFTM SEQ II LENG US-10-24 US-10-24
61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120	Query Match 100.0%; Score 663; DB 14; Length 132; Best Local Similarity 100.0%; Pred. No. 4.3e-66; Best Local Similarity 100.0%; Pred. No. 4.3e-66; Matches 132; Conservative 0; Mismatches 0; Indels 0; Gaps 0; 1 AKLETVTIGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60	WESULT 1 (Sequence 24, Application US/10243739) Publication No. US20030091593A1 GENERAL INFORMATION: APPLICANT: Bachmann, Martin F. APPLICANT: Storni, Taxio APPLICANT: Storni, Taxio APPLICANT: Lechner, Franziska TITLB OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of ITTLB OF INVENTION: Inmune Responses Induced by Virus Like Particles FILE REFERENCE: 1700.0210001 CURRENT APPLICATION NUMBER: US/10/243,739 CURRENT FILING DATE: 2002-09-16 PRIOR APPLICATION NUMBER: 60/318,967 PRIOR FILING DATE: 2001-09-14 NUMBER OF SEQ ID NOS: 73 SOFTWARE: Patentin version 3.1 SEQ ID NO 24 LENGTH: 132 TYPE: PRT ORGANISM: Bacteriophage Q-beta

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; ORGANISM: Bacteriophage Q-beta
US-10-244-065-24
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SEQ ID NO 24
LENGTH: 132
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Publication No. US20030157479A1
GENERAL INFORMATION:
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Best Local Similarity
                                                         APPLICANT: Jennings, Gary
APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/1802/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
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CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2002-04-22
PRIOR APPLICATION NUMBER: 60/318,994
PRIOR FILING DATE: 2001-09-14
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FILING DATE: 2002-01-18
APPLICATION NUMBER: US 60/331,045
FILING DATE: 2001-11-07
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||IDAIDQLNPAY 132
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Renhofa, Regina
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Lipowsky, Gerad
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; SOFTWARE: PatentIn Ver. 2; SEQ ID NO 256; LENGTH: 132; TYPE: PRT; ORGANISM: Qb 243
US-10-050-902-256
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SOFTWARE: PatentIn version
SEQ ID NO 24
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Best Local Similarity
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Best Local Similarity
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PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
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PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
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TITLE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0190004
CURRENT APPLICATION NUMBER: US/10/050,902
CURRENT FILING DATE: 2002-01-18
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                            121 LIDAIDQLNPAY 132
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                                                                         61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                        61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTPSFTQYSTDEERAFVRTELAALLASPL 120
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LIDAIDQLNPAY 132
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Tissot, Alain
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Sebbel, Peter
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US-10-050-898-256

Application US/10050898 o. US20030175711A1

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; LENGTH: 132
; TYPE: PRT
; ORGANISM: Qb 243
US-10-050-898-256
APPLICANT: Bachmann, Martin
APPLICANT: Maurer, Patrick
APPLICANT: Maurer, Patrick
APPLICANT: Pellicioli, Erica
APPLICANT: Renner, Wolfgang A.
TITLE OF INVENTION: Prion Protein Carrier-Conjugates
FILE REFERENCE: 1700.0290003
CURRENT APPLICATION NUMBER: US/10/346,190
CURRENT APPLICATION NUMBER: 60/396,590
PRIOR APPLICATION NUMBER: 60/396,590
PRIOR FILING DATE: 2002-07-18
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/389,898
PRIOR APPLICATION WHEER: 60/389,898
PRIOR FILING DATE: 2002-06-20
PRIOR FILING DATE: 2002-06-20
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FILE REFERENCE: 1700.0190005
CURRENT APPLICATION NUMBER: US/10/050,898
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR PPLICATION NUMBER: US 60/288,549
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR APPLICATION NUMBER: US 60/286,998
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-10-05
                                                                                                                                                                                                                                                                                                             Sequence 24, Application US/10346190 Publication No. US20030219459A1 GENERAL INFORMATION:
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PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 350
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 256
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Best Local 9
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APPLICANT:
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Tissot, Alain
Maurer, Patrick
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Sebbel, Peter
Piossek, Christine
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Ortmann, Rainer
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RESULT 7
US-10-465-811-15
; Sequence 15, Application US/10465811
; Publication No. US20040005338A1
; GENERAL INFORMATION:
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; TYPE: PRT
; ORGANISM: Qb :
US-10-346-190-24
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                                                                                                                                                                                                                                  Query Match 100.0%; Score 663; DB 15; Best Local Similarity 100.0%; Pred. No. 4.3e-66;
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SEQ ID NO 15
LENGTH: 132
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/465,811
CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: US 60/389,898
PRIOR FILING DATE: 2002-06-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: RENNER, WOLFGANG A
TITLE OF INVENTION: PACKAGED VIRUS-LIKE PARTICLES FOR USE AS ADJUVANTS:
TITLE OF INVENTION: METHOD OF PREPARATION AND USE
FILE REFERENCE: 1700.0290004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR FILING DATE: 2002-01-18
NUMBER OF SEQ ID NOS: 164
                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 131
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                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Bacteriophage Q-beta 243 mutant
                                                                                                                                                                                                                   132;
121
                      121 LIDAIDQLNPAY 132
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                                                                  61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                          61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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LIDAIDQLNPAY 132
                                                                                                                                           AKLETVTLGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                                                                                                                                                                                                   Conservative
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ilarity 100.0%;
Conservative (
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Pred. No. 4.3e-66;
D; Mismatches 0;
                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                     DB 15;
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RESULT 8
US-10-289-456-24
; Sequence 24, Application US/10289456

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CURRENT APPLICATION NUMBER: US/10/289,456
CURRENT FILING DATE: 2002-11-07
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/50,902
PRIOR PILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR PILING DATE: 2002-07-19
PRIOR FILING DATE: 2002-07-19
PRIOR FILING DATE: 2002-11-07
NUMBER OF SEQ ID NOS: 170
NUMBER OF SEQ ID NOS: 170
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US-10-622-064-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 7, Application US/10622064 Publication No. US20040059094A1
                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                            APPLICANT: Maurer, Patrik F
IITLE OF INVENTION: Hapten-Carrier Conjugates and Uses Thereof
FILE REFERENCE: 1700.0300001
CURRENT APPLICATION NUMBER: US/10/622,064
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,575
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APPLICANT: Maurer, Patrick
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EQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bachmann, Martin F APPLICANT: Maurer, Patrik F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Q-beta 243 mutant
                                                                                                                                                                                                                                 TYPE: PRT
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                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                       OTHER INFORMATION: Bacteriophage Q-beta 243 mutant
                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                        ENGTH: 132
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                                                                  132;
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132; Conserv
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spohn, Gunther
spohn, antigen
                                                                                       Similarity
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AKLETYTLGKIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRYTYSVSQPSRNKK 60
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                                                                  100.0%; Score 663; DB 15; ilarity 100.0%; Pred. No. 4.3e-66; Conservative 0; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/10/622,124
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,638
PRIOR FILING DATE: 2002-07-19
NUMBER OF SEQ ID NOS: 146
SOFTWARE: Patentin version 3.2
SEQ ID NO 18
LENGTH: 132
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US-10-622-124-18
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                                                           APPLICANT: Frey, Peter
TITLE OF INVENTION: Amyloid Beta 1-6 Antigen Ar
FILE REFERENCE: 1700.0350002
CURRENT APPLICATION NUMBER: US/10/622,087
CURRENT FILING DATE: 2003-07-18
CURRENT APPLICATION NUMBER: US 60/396,639
PRIOR APPLICATION NUMBER: US 60/470,432
PRIOR APPLICATION NUMBER: US 60/470,432
PRIOR FILING DATE: 2003-05-15
PRIOR FILING DATE: 2003-05-15
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Best Local Similarity
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APPLICANT: Fulurija, Alma
TITLE OF INVENTION: Ghrelin-Carrier Conjugates
SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 1700.0340001
                                                                                                                                                                                                                                                                                               APPLICANT: Bachmann, Martin F
APPLICANT: Tissot, Alain
APPLICANT: Ortmann, Rainer
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ORGANISM: Artificial Sequence
FEATURE:
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SOFTWARE: PatentIn version 3.2 EQ ID NO 18
                                         NUMBER OF SEQ ID NOS: 93
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Staufenbiel, Matthias
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Pred. No. 4.3e-66;
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US-10-243-739-26
US-10-243-739-26
; Sequence 26, Application US/10243739
; Publication No. US20030091593A1
; GENERAL INFORMATION:
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US-10-243-739-23
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APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Lechner, Franziska
TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
FILE REFERENCE: 1700.0210001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 23
LENGTH: 132
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Best Local Similarity
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Best Local Similarity
                APPLICANT: Bachmann, Martin F. APPLICANT: Storni, Tazio APPLICANT: Lechner, Franziska
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ORGANISM: Artificial Sequence
FEATURE:
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In vivo Activation of Antigen Presenting Cells for Enhancement
                    Franziska
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Pred. No. 3.4e-65;
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APPLICANT: Sciwarz, Katrin
APPLICANT: Meijerink, Edwin
APPLICANT: Meijerink, Edwin
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Pumpens, Paul
APPLICANT: Renhofa, Regina
ITILE OF INVENTION: Method of Immunostimulatory Sub
ITILE OF INVENTION: Method of Preparation and Use
FILIE REFERENCE: 1700.0220001
CURRENT APPLICATION NUMBER: US/10/244,065
CURRENT APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR FILING DATE: 2002-09-16
PRIOR FILING DATE: 2001-09-14
INUMBER OF SEQ ID NOSE: 73
SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
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SEQ ID NO 26
LENGTH: 132
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Publication No.
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
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Pred. No. 3.4e-65;
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APPLICANT: Meijerink, Edwin
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Pumpens, Paul
APPLICANT: Pumpens, Paul
APPLICANT: Renhofa, Regina
TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Particl
TITLE OF INVENTION: Method of Preparation and Use
FILE REFERENCE: 1700.0220001
CURRENT APPLICATION NUMBER: US/10/244,065
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/318,994
PRIOR APPLICATION NUMBER: 60/318,994
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 26
LENGTH: 132
TYPE: PRT
ORGANISM: Bacteriophage Q-beta
US-10-244-065-26
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US-10-244-065-26
US-10-244-065-26; Sequence 26, Application US/10244065; Publication No. US20030099668A1
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Tissot, Alain
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S01964
readthrough protein - phage SP
C;Species: phage SP
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004

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Nucleic Acids Res. 16, 6205-6221, 1988

A,Title: Analysis of the complete nucleotide sequence of A,Reference number: S01963; MUID:88289362; PMID:3399390
A,Accession: S01964
A,Status: preliminary
A,Molecule
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A;Note: the authors translated the readthrough stopcodon
C;Superfamily: phage GA coat protein
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C;Species: Listeria seeligeri
C;Date: 22-Nov-1993 #sequence_revision 01-Dec-1995 #text
C;Accession: S22340
R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete sequence of ilo f
A;Reference number: S22340; MUID:92182018; PMID:1543752
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R;Dhaese, P.; Vandekerckhove, J.S.; Van Montagu, Eur. J. Bicchem. 94, 375-386, 1979
A;Title: The primary structure of the coat protei A;Reference number: A04225; MUID:79148387; PMID:1A;Accession: A04225
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Pred. No. 0.00035;
4; Mismatches 58;
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C;Accession: T00646

R;Federspiel, N.A.; Palm, C.J.; Conway, A.B.; Kurt; Vysotskaia, V.S.; Yu, G.; Ecker, J.; Theologis, submitted to the EMBL Data Library, February 1998 A;Reference number: Z14197
A;Accession: T00646
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A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-530 <HAA>
A;Cross-references: UNIPROT:P31830; EMBL:X60462;
A;Note: the authors translated the codon GCC for C;Superfamily: dipeptide transport protein
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A;Introns: 14/3; 281/3; 428/3; 448/2;
A;Introns: 14/3; 281/3; 428/3; 448/2;
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A;Reference number: S24230;
A;Accession: S24231
A;Status: nucleic acid seque
A;Molecule type: DNA
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R;Rasmussen, O.F.; Beck, T.; Olsen, Infect. Immun. 59, 3945-3951, 191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Gene: ATSP:F3I6.9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical
                                                                                                         A;Title: Listeria monocytogenes isolates can be classified into A;Reference number: S24230; MUID:92040062; PMID:1937753
                                                                                                                                                                                                                                                                           C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995
                                                                                                                                                                                                                                                                                                               C;Species: Listeria monocytogenes A;Variety: strain 12067
                                                                                                                                                                                                                                                                                                                                                                           listeriolysin precursor - Listeria monocytogenes (strain 12067)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ypothetical protein F3I6.9 - Arabidopsis thaliana;Species: Arabidopsis thaliana (mouse-ear cress);Date: 01-Feb-1999 #sequence_revision 01-Feb-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11 IGKDGKOTIVLNPRG---VNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKN-YKVQV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               34;
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Similarity 27.0%;
31; Conservative 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13.7%; Score 90.5; DB
25.0%; Pred. No. 0.56;
cive 22; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 88.5; DB Pred. No. 0.89;
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(lipoamide)
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residue 287
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A;Residues: 1-529 <RAS>
A;Cross-references: UNII
                                                                                 A;Residues: 413-480 <MEZ-
A;Cross-references: GB:M29171
R;Michel, E.; Reich, K.A.; Favier, R.; Berche, P.; Cossart, P.
Mol. Microbiol. 4, 2167-2178, 1990
Mol. Microbiol. 4, 2167-2178, 1990
A;Title: Attenuated mutants of the intracellular bacterium Listeria monocytogenes
A;Reference number: S12400; MUID:91211627; PMID:1965218
                                                                                                                                                                                                                                                                                                                                    A;Cross-references: EMBL:X15127; NID:g44106; PIDN:ChA33223.1; PID:g44107 A;Experimental source: strain EGD A;Note: this sequence is derived from a weakly hemolytic strain, serotype R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P. Infect. Immun. 55, 3225-3227, 1987 A;Title: Identification of the structural gene encoding the SH-activated h A;Reference number: A47606; MUID:88057627; PMID:2824384 A;Accession: A47606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Listeria monocytogenes
C;Date: 21-Oct-1992 #sequence revision 21-Oct-1992 #text_change 09-Jul-2004
C;Accession: A43505; 805306; A47606; S12400; A61079
R;Mengaud, J.; Vicente, M.F.; Chenevert, J.; Pereira, J.M.; Geoffroy, C.; Gicquel-Sanzey Infect. Immun. 56, 766-772, 1988
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C;Keywords: virulence factor
F;1-25/Domain: signal sequence #status predicted
F;26-529/Product: listeriolysin #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPROT:P13128; GB:M24199; NID:g149652; PIDN:AAA03018.1; PID:g149653; A;Note: this sequence is derived from a strongly hemolytic strain, serotype 1/2c R;Domann, E.; Chakraborty, T.
Nucleic Acids Res. 17, 6406, 1989
A;Title: Nucleotide sequence of the listeriolysin gene from a Listeria monocytogenes ser A;Reference number: S05306; MUID:89366684; PMID:2505236
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                               A;Molecule type: DNA
A;Residues: 483-493 <MIC>
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;Experimental source: strain 12067, serotype 4b
;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1991
     Experimental source:
                                                                                                                                                                                                                                                                                                          Status: preliminary
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;Residues: 1-529 <DOM>
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Pred. No. 2.
serotype 1/2c
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                                                                                    T31660
hypothetical protein COS41.6 - sea squirt (Ciona intestinalis)
C;Species: Ciona intestinalis
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_chang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, B.; Maitournam, ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; Voss, H.; Wa;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Accession: AC1100
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C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 09-Jul-2004
C;Accession: AC1100
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                                   R;Bird, A.P.; Clark, V.; Jones, S.J.; Leitgeb, submitted to the EMBL Data Library, December 19
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C;Superfami
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A; Residues: 1-529 <GLA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloec.; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.; Fsihi,
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                    A; Reference number:
                                                                            C; Accession: T31660
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A; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               listeriolysin O precursor [imported] - Listeria monocytogenes (strain EGD-e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13 KDGKQTLVL------NPRGVNPTNGVASLSQAGA-------VPALEKRV 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30;
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22.7%; Pred. No. 2.1;
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22.7%;
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                                                       Dobson, R.; Tweedie, S
                                                                                           #text_change 09-Jul-2004
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Wehland,
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A;Status:

preliminary; translated from

GB/EMBL/DDBJ

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A;Cross-
A;Note: 1
R;Kreft,
CYC1/CYP3 transcription activator - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein 19672.1; protein YLR256w; regulatory protein CYP:
C;Species: Saccharomyces cerevisiae
C;Date: 30-Sep-1991 #sequence revision 23-Feb-1996 #text_change 12-Nov-1999
C;Accession: S59400; A31312; S15447; S05804; S15446
R;Johnson, D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Accession: S22341; S36683
R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete sequence of ilo from A;Reference number: S22340; MUID:92182018; PMID:1543752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Listeria ivanovii
C;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 09-Jul-2004
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A; Residues: 1-432 <BIR>
A; Cross references: UNIPROT: P91584;
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A;Residues: 1-319,'T',321-528 <KRE>
A;Cross-references: EMBL:X60461
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A;Molecule type: DNA
A;Residues: 1-528 <HAS>
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Note: the authors translated the c
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                                                                                                                                                                                                                                                                                                           TVSVSQPSRNRKNYKVQVKIQNPTACTANGSCDPSVTR-----QKYADVTFSFTQYSTD 102
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28.3%;
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Pred. No. 5.3;
24; Mismatches
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codon ACA for
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                                                                              protein CYP1;
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                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                            regulatory
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submitted to the EMBI
A;Description: The se
A;Reference number: S
A;Accession: S59400
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A;Map position: 12R
C;Superfamily: regulatory protein HAP1; GAL4 zinc binuclear clust
C;Keywords: DNA binding; heme binding; transcription regulation;
E;1-148/Domain: DNA binding #status predicted <DNA>
F;59-98/Domain: GAL4 zinc binuclear cluster homology <GAL4>
F;64-84/Region: zinc finger CCC motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Rosidues: 1-144,'I',146-322,'R',324-454,'N',456-507,'M',509-586,'K',588-882,'N',
A; Residues: 1-144,'I',146-322,'R',324-454,'N',456-507,'M',509-586,'K',588-882,'N',
A; RCCOSS-references: EMBL.J03152; NID:g171645; PIDN:AAA34662:1; PID:g171646
R; CCCUSOt, F:; Verdiere, J:; Gaisne, M:; Slonimski, P.P.
J. MOl. Biol. 204, 263-276, 1988
A; Title: CYPI (HAPI) regulator of oxygen-dependent gene expression in yeast. I. Ov
A; Reference number: S15447; MUID:89125585; PMID:2851658
A; Accession: S15447; MUID:89125585; PMID:2851658
                                                                                                                                                                                                                                                                                                                                                                                                                                  F;177-189/Region: glutamine-rich
F;245-445/Domain: heme binding #status predicted <HEM>
F;245-445/Domain: heme binding #status predicted <FEY259-304,323-328,347-352,373-378,389-394,415-420/Region:
F;1308-1481/Domain: activation element #status predicted F;1308-1481/Region: acidic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: DNA
A;Residues: 1-62,'R',64-1305,'Y',1306-1470,'LVDFYRADFPIWE'
A;Cross-references: EMBL:X13793
A;Note: the sequence is from mutant CYP1-18
C;Genetics:
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A;Cross-references: SGD:S0004246; MI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cell 56, 291-301, 1989
A;Title: Functional dissection and sequence of yeast HAP1
A;Reference number: A31312; MUID:89106221; PMID:2643482
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R; Pfeifer, K.; Kim, K.: Cell 56, 291-301, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Residues: I-1502 <JOH>
A;Cross-references: EMBL:U20865; NID:g662330;
A;Experimental source: strain S288C (AB972)
망
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Best Local S
Matches 35
                                                                                                         1338 LPDPSRDEAFDNSIKOMTPMTSAFMNANTTIPSSTLNGNMNMNGAGTANTDTSANGSALS
                                                                                                                                                                                                                     1278 PRGISPKPSNGLSSVQPLLSSFSMNQLNGGTIPTVPSLTNITSQMGALPSLDRITTNQIN
                                                                                                                                                               54 QPSRNR-KNYKVQVKIQNP-----
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                                                                                                                                                                                                                                                                                                                                                        Similarity
                                             SVTRQKYADV-TFSFTQYSTD-EERAFVRTELAALLASP-LLIDAIDQLN 129
TLTSPQGSDLASNSATQYKPDLEDFLMQNSNFNGLMINPSSLVEVVGGYN 1447
                                                                                                                                                                                                                                                                        PRGVN--PTNGVASL-----
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The sequence of S. cerevisiae cosmid 9672.
mber: S59386
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                                                                                                                                                                                                                                                                                                                                                  12.1%;
                                                                                                                                                                                                                                                                                                                              30;
                                                                                                                                                                                                                                                                                                                           Score 80.5; D
Pred. No. 18;
30; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PIDN: AAB67387.1;
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                                                                                                                                                                                                                                                                           ---SQAGAVPALEKRVTVSVS
                                                                                                                                                                                                                                                                                                                                                                                 Length 1502;
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                                                                                                                                                                                                                                                                                                                                 63;
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884-959

A; Reference number: A; Accession: C98221 A; Status: prelimina: R;Goodner, B.; Hinkle, G.; Gattung, S.; Miller, N.; Blancha A.; Liu, F.; Wollam, C.; Allinger, M.; Doughty, D.; Scott, Science 294, 2323-2328, 2001 C;Date: 22-Oct-2001 #sequence_revision
C;Accession: C98221 hypothetical protein AGR_L_1428 [imported] - Agrobacterium tumefaciens (strain C;Species: Agrobacterium tumefaciens A; Title: Genome Sequence

A; Molecule type: DNA A; Residues: 1-136 < KUR> UNIPROT: Q8U8F3; GB: AE007870; PIDN:AAK89293.1; PID:g15159127; GSPDB:GN

uence of the Plant Pathogen and Biotechnology A97359; MUID:21608551; PMID:11743194

Agent Agrobacterium tume

B.; Goldn Markelz,

Goldman, kelz, B.;

C58,

Cerec

22-Oct-2001 #text_change 09-Jul-2004 Miller, N.; Blanchard, M.; Qurollo, Doughty, D.; Scott, C.; Lappas, C.;

preliminary

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R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, M.B deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, G.; il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barrell, Nature 413, 523-527, 2001
                           A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-282 <KUR>
                                                                                                  A;Title: Genome sequence of Yersinia pestis, the causative agent of plague A;Reference number: AB0001; MUID:21470413; PMID:11586360
A;Accession: AI0186
                                                                                                                                                                                                                                                                                                                                                probable iron-siderophore transport system, ATP-binding component YPO1533 [imported] -
C;Species: Yersinia pestis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT:Q8U8F3; GB:AE008689; PIDN:AAL44939.1; PID:g17742593; GSPDB.
A;Experimental source: strain C58 (Dupont)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58 A;Reference number: AB2577; MUID:21608550; PMID:11743193 A;Accession: AE3065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , Karp, P.; Romero, P.; Zhang, S.
Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Genetics:
A;Gene: AGR_L_1428
A;Cross-references: UNIPROT:Q8ZFZ0; GB:AL590842; PIDN:CAC90356.1; PID:g15979576; GSPDB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   conserved hypothetical protein Atu4139 [imported] - Agrobacterium tumefaciens
C;Species: Agrobacterium tumefaciens
                                                                                                                                                                                                                                                                                       Species: Yersinia pestis
Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Jul-2004;
Accession: A10186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gene: Atu4139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 09-Jul-2004
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11 Similarity 25.4%;
30; Conservative 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        60 KNYKVQVKIQNPTACTA----NGSCDPSVTRQKYADVTFSF-TQYSTDEERAFVRTE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11 IGKDGKQTLVLNPRGVNPTNGVASLSQAGAV------PALEKRVTVSVSQPSRNR 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60 KNYKVQVKIQNPTACTA----NGSCDPSVTRQKYADVTFSF-TQYSTDEERAFVRTE 111
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30; Conserv
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25.4%; Pred. No. 1.4;
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Pred. No. 1.
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Search completed: January Job time: 9.03224 secs

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A;Cross-references: UNIPROT:Q99QU5; GB:BA000018; PID:g13702654; PIDN:BAB43794.1; GSPDB:Gh
A;Experimental source: strain N315
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.; ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimizu, C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K. Lancet 357, 1225-1240, 2001

A;Title: Whole genome sequencing of meticillin-resistant Stapylococcus aureus. A;Reference number: A89758; MUID:21311952; PMID:11418146

A;Accession: H90078
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C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 09-Jul-2004
C;Accession: H90078
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56 LHFKTEVQSSLPFKLTGDVSCNHAPKRQQMIDFIVATAALDIELLRSLVTTE 107
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"Evolution of phage with chemically ambiguous proteomes.";
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01-OCT-2002 (TrEMBLrel. 22, Last sequence
05-JUL-2004 (TrEMBLrel. 27, Last annotati
Al read-through protein (Al protein).
Bacteriophage Q-beta.
Viruses; ssRNA positive-strand viruses, nc
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PubMed=14667253;

Bacher J.M., Bull J.J., El

"Evolution of phage with c

"BMC Evol. Biol. 3:24-24(20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; IEA.
InterPro; IPR002703; Levi coat.
InterPro; IPR000504; RNA_Tec_mot.
Pfam; PF01819; Levi coat; 1.
PROSITE; PS00030; RRM_RNP_1; UNKNOWN 1.
SEQUENCE 329 AA; 36134 MW; E0F30215A8C197CB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE=94109687; PubMed=7506687;
Kozlovska T.M., Cielens I., Dreilinna D.,
Ose V., Pumpens P.;
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EMBL; AY099114; AAM33127.1;
EMBL; M99039; AAA16663.1; -.
HSSP; P03615; 1QBE.
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Q9TOR9;
01-MAY-2000
                    GO; GO:0019028; C:viral capsid; IEA. GO; GO:0005198; F:structural molecule InterPro; IPR002703; Levi coat. Pfam; PF01819; Levi coat; 1. Coat protein.
                                                                                                                        Beekwilder M.J., Nieuwenhuizen R., van Duin J. Submitted (APR-1998) to the EMBL/GenBank/DDBJ EMBL; AF059242; AAC14699.1; -... HSSP; P03615; 1QBB.
                                                                                                                                                                                                                                                         Beekwilder J., Nieuwenhuizen R., Po
"Secondary structure model for the
Control of A-protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=95239761; PubMed=7723040; Beekwilder M.J., Nieuwenhuizen R., "Secondary structure model for the RNA phage Q beta."; J. Mol. Biol. 247:903-917(1995).
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MEDLINE=94109687; PubMed=7506687;
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Viruses; ssRNA positive-strand
Allolevivirus
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Ose V., Pumpens P.;
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Best Local Similarity
Matches 110; Conserv
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InterPro; IPR002504; RNA_rec_mot.
Pfam; PF01819; Levi_coat; 1.—
PROSITE; PS00030; RRM RNP 1; UNKNOWN
SEQUENCE 329 AA; 35880 MW; 372BI
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HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule
                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=95239761; PubMed=7723040; Beekwilder M.J., Nieuwenhuizen R. "Secondary structure model for th RNA phage Q beta."; J. Mol. Biol. 247:903-917(1995).
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01-AUG-1998 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
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Allolevivirus.
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Best Local Similarity
Matches 106; Conserv
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O1-AUG-1998 (TrEMBLrel. 07, Creat O1-AUG-1998 (TrEMBLrel. 07, Last O1-MAR-2004 (TrEMBLrel. 26, Last A1-protein.
Bacteriophage M11.
Viruses; ssRNA positive-strand vi Allolevivirus.
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GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
Coat protein.
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HSSP; P03615; 1QBE.
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      MEDLINE=95239761;
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                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content use by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
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HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity;
InterPro; IPR002703; Levi_coat.
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.

PROSITE; PS00030; RRM RNP 1; UNKNOWN 1.

SEQUENCE 329 AA; 35893 WW; 3E33CD821EE625F4
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    EMBL;
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                                                                                                                entities requires a license agreement (some send an email to license@isb-sib.ch)
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                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=88289362; PubMed=3399390;
Hirashima A., Hirose T., Inayama S., Inokuchi Y., Jacobson "Analysis of the complete nucleotide sequence of the group coliphage Sp.";
Nucleic Acids Res. 16:6205-6221(1988).
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01-MAR-1989 (Rel. 10, Last sequence update)
29-MAR-2004 (Rel. 43, Last annotation updat
Readthrough protein Al [Contains: Coat prot
Bacteriophage SP.
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P09677;
                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collabora between the Swiss Institute of Bioinformatics and the EMBL outstati the European Bioinformatics Institute. There are no restrictions on use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commer entities requires a license agreement (See http://www.isb-sib.ch/annou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Viruses; serna positive-strand viruses, no DNA stage; allolevivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR002703; Levi_coat. Pfam; PF01819; Levi_coat; 1. Coat protein; RNA-binding. SEQUENCE 132 AA; 14129 MW;
                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                  InterPro; IPR002703; Levi coat. Pfam; PF01819; Levi coat; 1.
                                                                                                                                                                                                                              HSSP; P03615; 1QBE
                                                                                                                                                                                                                                                        EMBL; X07489; CAB37299.1; -.
                                                                                                                                                                                                                                                                                   or send an email to license@isb-sib.ch).
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NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
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                                        AKLNQVTLSKIGKNGDQTLTLTPRGVNPTNGVASLSEAGAVPALEKRVTVSVAQPSRNRK
                                                                                                                                               331 AA;
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79.5%;
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79.5%; Pred.
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                                                                                            Score 537.5; DB 1; Length Pred. No. 9.2e-44; 2; Mismatches 14; Indels
                                                                                                                                             Coat protein.
Readthrough protein A1.
, 4A6642E4B52C6582 CRC64;
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No. 3.2e-44;
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nt protein].
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01-AUG-1998
01-MAR-2004
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Q9TOR8;
01-MAY-2000
                                                                                                                                            064310;
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HSSP; P036.15; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity;
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
Enterobacteria phage NL95. Viruses; ssRNA positive-strand viruses,
                                           A1-protein.
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"Secondary structure model for the first three
Control of A.protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
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MEDLINE=96190948; PubMed=8609616;
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MEDLINE=95239761; PubMed=7723040;
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Biol. 247:
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GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; I
GO; GO:0005198; F:structural molecule activity; I
InterPro; IPR002703; Levi_coat.
InterPro; IPR000504; RNA_rec_mot.
Pfam; PF01819; Levi_coat; 1.
PROSITE; PS00030; RRM_RNP 1; UNKNOWN 1.
SROUENCE 330 AA; 36175 MW; 961E55F408334410
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COAT_E
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                                                                                                                                                                                                                                                                                                                                            Bacteriophage PRR1.
Viruses; ssRNA positive-strand viruses,
unclassified Leviviridae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Beekwilder M.J., Nieuwenhuiz
Submitted (APR-1998) to the
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InterPro; IPR002703; Levi coat.

Pfam; PF01819; Levi coat; 1.

Coat protein; Direct protein sequencing; RNA-binding.
SEQUENCE 131 AA; 14535 MW; E7E639E1E50FC612 CRC64;
                                                                                             Eur. J. Biochem. 94:375-386(1979)
-!- FUNCTION: Forms the phage she PIR, A04225, VCBPP1.
                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JUL-1986 (Rel. 01,
21-JUL-1986 (Rel. 01,
05-JUL-2004 (Rel. 44,
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                                                                                                                                                                        Dhaese P., Vandekerckhove J., "The primary structure of the bacteriophage PRR1.";
                                                                                                                                                                                                                                  SEQUENCE.

MEDLINE=79148387; PubMed=107028;
                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=12024;
                                                                                                                                                                                                                                                                                                                                                                                                                     Coat protein.
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21-JUL-1986
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Last
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Matches 42
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01-JUL-1993 (Rel.
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Seeligeriolysin F
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Q8VDC2;
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LISSE
TACY LISSE
P31830;
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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Mamm. Genome 13:646-655(2002).
EMBL; AJ428064; CAD20986.1; -.
Listeria seeligeri.
Bacteria, Firmicutes,
                                          Name=1so;
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MEDLINE-22350095; PubMed-12461651;
Kiss H., Darai E., Kiss C., Kost-A
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Rodentia;
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Pred. No. 0.0027
4; Mismatches
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Best Local Similarity
Matches 34; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Listeriolysin genes: complete sequence of ilo from Listeria ivanovii and of 1so from Listeria seeligeri." Biochim. Biophys. Acta 1130:81-84(1992).

-I- FUNCTION: Sulfhydryl-activated toxin. Is able to lyse cholesterol containing membranes. Can be reversibly inactivated by oxidation. Cholesterol 1s the receptor for the binding of these toxins to eukaryotic cell membranes.

-i- SIMILARITY: Belongs to the thiol-activated cytolysin family.
                                                                                                                                                                                                                                                                                                                                                                                                             CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; X60462; CAA42996.1; -.

PIR; S23340; S2340; S2340.

HSSP; P1995; 1PFO.

InterPro; IPR001869; Thiol_cytolysin.

Pfam; PF01289; Thiol_cytolysin; 1.

PRINTS; PR01400; TACYTOLYSIN.

PRODOm; PD007062; Thiol_cytolysin; 1.

PROSTTE; PS00481; THIOL_CYTOLYSINS; 1.

CYtolysis; Hemolysis; Lipid-binding; Signal; Toxin.

SIGNAL

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
STRAIN=SLCC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=1640;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Haas A., Dumbsky M., Kreft J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=92182018; PubMed=1543752;
                                                                             209
                                                                                                                                                    154 TLSVDLPGMTKKDNKIFVK--NPTKSNVNNAVNTLVERWNDKYSKAYPNINAKIDYS--- 208
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                                                                                                                                                                                                                                                                       13
                                                                                                             99 YSTDEBRAFVRTELAA 114
                                                                                                                                                                          TVSVSQPSRNRKNYKVQVKIQNFTACTANGSCDPSVTR--QKY------ADVTFSFTQ 98
                                                                             ----DÉMÁYSESQLIÁ 220
                                                                                                                                                                                                                                                                       KDGKQTLVL--NPRGVNPT-----NGVASLSQAGA------VPALEKRV 48
                                                                                                                                                                                                                                 KDGSBYIVVEKKKKGINQNNADISVINAISSLTYPGALVKANRELVENQPNVLPVKRDSL 153
                                                                                                                                                                                                                                                                                                                                                                                   26 530 Seeligeriolysin.
485 485 Binding to cholesterol (By similarity).
530 AA; 59181 MW; 416F7A4DD2029866 CRC64;
                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                          13.7%; Score 90.5; DB 1; Length 530; 25.0%; Pred. No. 4.1; tive 22; Mismatches 37; Indels 4.
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GenCore version (c) 1993 - 2005

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                geneseqp2002s:*
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Result

Query

Renner WA, Piossek C;

Bachmann M,

Tissot A,

Maurer P,

Lechner

'n

Sebbel P, (CYTO-) CYTOS BIOTECHNOLOGY AG.

WPI; 2002-627351/67.

This invention relates to a novel ordered and repetitive antigen array used in the production of vaccines for infectious diseases. The invention also discloses a composition comprising a non-natural molecular scaffold comprising a core particle selected from a core particle of a non-natural origin and a core particle of natural origin and an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond. Also disclosed is an antigen or antigenic determinant with at least one second attachment

Claim 18; Page 144; 441pp; English.

diseases. Molecular

antigen array used in the

production

of vaccines

for infectious

45	44	43	42	41	40	39	38	37	36	35	34		32	31	30	29	28	27	0
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Abu09692	Abu09690	Abu09686	Abr56453	Abr56439	Abr56456	Abg80545	Abg80631	Abg80629	Abg94319	Abg94317	Abg94233	Adk52196	Adj67162	Adj36316	Adk17147	Adj82059	Add24133	Abr44558	Audososs
Bacteriop	Bacteriop	Bacteriop	Bacteriop				Bacteriop	Bacteri		pQB243 pr	Bacteriop		Phage Qbe	Bacteriop		Protein	Bacteriop		

ALIGNMENTS

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RESULT 1
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ABG9416
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04-MAY-2001; 2001US-0288549P.
05-OCT-2001; 2001US-032699BP.
07-NOV-2001; 2001US-0331045P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory; cytostatic; antiviral; antidiabetic; hypoglycaemic; antigen array; vaccine; infectious disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unidentified bacteriophage.
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10-DEC-2002
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                                              19-JAN-2001;
04-MAY-2001;
05-OCT-2001;
07-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Molecular antigen array; vaccine; antigen; antimicrobial; molecular scaffold; amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgB-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus erythematosus; osteoporosis; inflammatory immune disease; myasthenia gravis; multiple sclerosis; immunoproliferative disease; myasthenia gravis; multiple sclerosis; angioimmunoproliferative disease; impabadenopathy; Alzheimer's disease; angioimmunoproliferative lymphadenopathy; immunoblastive lymphadenopathy; rehematoid arthritis; diabetes; infectious disease; factor Xa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              capable of association through at least one non-peptide bond to the first attachment site and where the antigen or antigenic determinant and the scaffold interact through the association to form an ordered and repetitive antigen array. The invention also comprises a coat protein capable of forming a capsid which comprises mutant Obeta coat proteins having an amino acid sequence selected from five amino acid sequences fully defined in the specification. The compounds of the invention may have antimicrobial, antiallergic, immunomodulatory, cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiviral, antidiabetic, or hypoglycaemic activities and may be used in immunisation and as a vaccine. The present sequence represents a protein sequence used to create the compositions of the invention. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (Abeta1-42) or its fragment and where the second attachment site is selected from an attachment site not naturally occurring with the antigen or antigenic determinant and an attachment site naturally occurring with the antigen or antigenic determinant, where the second attachment site is capable of association through at least one non-peptide bond to the first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     enterokinase;
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                                              ; 2001US-0262379P.
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Comparising (i) a core particle selected from: (i) a core particle of a non-natural origin; and (2) a core particle of natural origin; and (2) a core particle of natural conginism and (3) a core particle of natural conginism and (2) a core particle of natural conginism; and (2) a core particle by at least one covalent bond; (b) an antigen or antigenic determinant with at least conginism site is connected to the core particle by at least conginism site is connected to the core particle by at least conginism site is connected to antigenic determinant with at least conginism site is selected from: (i) an attachment site not naturally occurring with the antigen or antigenic determinant; and (ii) an convenient site naturally occurring with the antigen or antigenic determinant; and the sociation of through the association to form an ordered and repetitive antigen array. The composition is used in conformation and as a vaccine for diseases such as influenza, graft or spiratory distress syndrome (ARDS), Crohn's disease, allergic asthma, conspiratory distress syndrome (ARDS), Crohn's disease, allergic asthma, consequence of systemic lupus erythematosus, inflammatory immune diseases, myasthenia conspiratory immune diseases, myasthenia conspiratory immune diseases, myasthenia consequence site (enteroxinase or factor Xa) and a Cysteine-containing N-cromanial linker peptide which serves as the attachment point to a containing site (enteroxinase or factor Xa) and a Cysteine-containing N-cromanial linker peptide which serves as the attachment point to a contain site is a contained to bacterial protein (the saffold protein). The containing N-cromanial linker peptide which serves as the attachment point to a contain site is a contained to the containing N-cromanial site is a contained to the containing N-cromanial site is a contained to the containing N-cromanial site is a contained to the co
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                                                                                                                                     virus like particle or bacterial protein (the scaffold present sequence is bacterial protein or peptide which modified antigen to form the molecular antigen array
                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a composition comprising: (a) a non-natural molecular scaffold comprising: (i) a core particle selected from: (1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Molecular antigen array used in the production of vaccines for infectious
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ORTMANN R.
LUEOEND R.
STAUFENBIEL M.
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                       Similarity
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Bachmann M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 144; 418pp; English
                                                                                          AA;
100.0%; Score 664; DB 5;
100.0%; Pred. No. 1.1e-70;
tive 0; Mismatches 0;
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, Tissot A,
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Piossek C;
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                                                                                                                                NYKVQVKI QNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERA FVRTELAALLASPL
                                                                                                                                                              AKLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
                                                                                    LIDAIDQLNPAY 132
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Sequence
132 AA;
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Bacteriophage Q-beta coat protein mutant SEQ ID NO:23.

28-JUL-2003

(first entry)

Antigen presenting cell; APC; immune response; virus like particle; VLP; cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory; antithyroid; antiidiabetic; neuroprotective; nootropic; osteopathic; antitheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis; allahemer's disease; osteoporosis; rheumatoid arthritis; inflammatory autoimmune disease.

Synthetic Bacteriophage Qbeta.

WO2003024480-A2

16-SEP-2002; 2002WO-IB004252.

14-SEP-2001; 2001US-0318967P

(CYTO-) CYTOS BIOTECHNOLOGY AG

H Lechner F;

WPI; 2003-363095/34.

A composition, useful for enhancing an immune response against an antigen or a virus-like particle, enhancing anti-viral protection in an animal, or immunizing or treating tumors or infectious diseases, e.g. viral

Disclosure; Page 179-180; 243pp; English

CC The present invention describes a composition (C) for enhancing an immune CC response against an antigen or a virus-like particle (VLP) bound to at least one antigen, or a CC Comprises a virus-like particle (VLP) bound to at least one antigen, or a CC VLP capable of being recognised by the immune system of the animal. Also CC comprising (1) enhancing an immune response against an antigen or a VLP CC in an animal comprising introducing (C) into the animal; (2) vaccines CC comprising (C) together with a pharmaceutical diluent, carrier or CC exciptent; (3) immunistage or treating an animal comprising administering the vaccine; and (4) enhancing anti-viral CC protection in an animal comprising introducing; and (7) enhancing anti-viral CC thas cytostatic, virucide, antibacterial, antiparasitic, fungicide, CC antitleysoid, antidabetic, neuroprotective, notropic, osteopathic, CC antitleysoid, antidabetic, neuroprotective, notropic, osteopathic, CC antitleysoid, antidatheritritic activities. (C) or the vaccines can be used for enhancing anti-viral protection in an animal, or immunising or treating tumours and infectious diseases such as viral, bacterial, CC parasitic or fungal infections. The vaccine compositions are also useful cfor preventing or treating allergies, drug addiction, graft-versus-host CC disease, Crohn's disease, osteoporosis, rheumatoid arthritis, or inflammatory autoimmune disease, osteoporosis, rheumatoid arthritis, or inflammatory cromoune disease, osteoporosis, rheumatoid arthritis, or inflammatory autoimmune disease. ACC69838 to ACC69852 and ABR56401 to ABR56509

Query Match Best Local Similarity Matches Conservative 100.0%; 0; Score 664; DB 6; Pred. No. 1.1e-70; 0; Mismatches 0; Length 132; Indels ٥, Gaps

AKLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60

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Query Match Best Local Similarity

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664; DB 6; No. 1.1e-70;

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                               The invention describes an angiotensin peptide moiety carrier conjugate comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kidney failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta coat protein used in the preparation of the vaccine conjugates of the invention
                                                                                                                                                                                                                                                                                                                    New angiotensin peptide moiety carrier conjugate comprising a carrier and an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-IB000156.
19-JUL-2002; 2002US-0396637P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteriophage Qbeta; coat protein; hypotensive; cerebroprotective; cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction; congestive heart failure; kidney failure; retinal haemorrhage; mutant;
                                                                                                                                                                                                                                                                  Claim 20; Page 93-94; 97pp; English.
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Pumpens P,
introducing (C) into the animal; (2) producing (C) for immune response in an animal; (3) vaccines comprising pharmaceutical diluent, carrier or excipient; and (4)
                                                                                                             The present invention describes a composition (C) for enhancing an imm response in an animal. (C) comprises a virus-like particle (VLP), and immunostimulatory substance. The immunostimulatory substance is bound the VLP. Also described: (1) enhancing an immune response in an animal
                                                                                                                                                                                                                                                                                                                                                                                                                New compositions comprising immunostimulatory substances packaged virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, turistics in the composition of the compositi
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                                                                                                                                                                                                                                                                                                              Disclosure; Page 259; 322pp; English.
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22-APR-2002;
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MEIJERINK E.

LIPOWSKY G.

PUMPENS P.

CIELENS I.

RENHOFA R.
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Cielens :
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2002US-0374145P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schwarz K, I
Renhofa R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Meijerink E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bachmann
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                               lke particle (VLP), and an cory substance is bound to be response in an animal by (C) for enhancing an rising (C) together with a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cc treating an animal by: (a) administering the vaccine to the animal; (b) priming a T cell response in the animal by administering the vaccine; or (c) boosting a T cell response in the animal by administering the vaccine; or (c) consting a T cell response in the animal by administering the vaccine. (C) has immunostimulant, cytostatic, antiallergic, virucide and contibacterial activities. (I) can be used in vaccines for enhancing an climmune response in an animal, particularly a mammal or human. T cell considered in vaccines for enhancing an considerable, (C) is useful for enhancing a B cell response, a T cell comprising (C) can also be used for immunising or treating an animal, ce.g. numans, sheep, horses, cattle, pigs, dogs, cats, birds, ce.g. humans, sheep, horses, cattle, pigs, dogs, cats, birds, considered constitution or synhilis. Acceptable infections (e.g. influenza, hepatitis, measles or chicken pox), or bacterial infections (e.g. tuberculosis, pneumonia or syphilis). Accepto to Accepts for the present formation of the present formation.
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Best Local S
Matches 132
                                                                                                                             18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-1B000166.
08-JUL-2002; 2002US-0397259.
18-JUL-2002; 2002US-0396590P.
                                                                                                                                                                                                                                                                                                                                                                     vaccine composition; virus-like particle; core particle; first attachment site; antigen; antigenic determinant; prion PrP; PrP peptide; vaccine; neuroprotective; antiinflammatory; prion disease; Bovine Spongiform Encephalopathy; BSE;
                                                                Bachmann
                                                                                                                                                                                                             17-JAN-2003; 2003WO-EP000460.
                                                                                                                                                                                                                                           24-JUL-2003
                                                                                                                                                                                                                                                                          WO2003059386-A2
                                                                                                                                                                                                                                                                                                          Bacteriophage Qbeta
                                                                                                                                                                                                                                                                                                                                                          Creutzfeldt-Jakob Disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteriophage Qbeta coat protein mutant Qbeta-240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    represent sequences used in the exemplification of the present invention
                                 2003-598483/56.
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                                                                                              CYTOS BIOTECHNOLOGY
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                                                                Maurer P,
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                                                                Pellicioli E,
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Pred. No. 1.1e-70;
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A vaccine composition for preventing

or treating prion diseases

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Best Local :
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like or a core particle with at least one first attachment site and at
least one antigen or antigenic determinant that is a prion protein (PrP)
or its dimer, or a PrP peptide, the antigen or antigenic determinant
being bound to the virus-like or core particle. The vaccine of the
invention may have neuroprotective or antiinflammatory activity. The
composition is useful as a medicament or in manufacturing a medicament
for the treatment or prevention of prion diseases. The prion diseases may
include Bovine Spongiform Encephalopathy (BSE) or Creutzfeldt-Jakob
Disease. The present sequence is the amino acid sequence of a mutant coat
protein from bacteriophage Obeta which may be used during the creation of
the vaccine composition of the invention.
New compositions comprising a core particle and at least one antigen or antigenic determinant, useful for as a vaccine for therapy or prophylaxis of bone diseases, particularly mammalian encephalopathies.
                                                                                                                                                                                                                                                                                               07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050992.
21-JAN-2002; 2002WO-1B000166.
19-JUL-2002; 2002US-0396635P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   osteopathic; bone disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-NOV-2002; 2002WO-EP012449.
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                                                                                                                          WPI; 2003-441430/41.
                                                                                                                                                                                                                                              (CYTO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein for RANKL antigen array to treat bone disease
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                                                                                                                                                                                 ' Maurer
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ilarity 100.0%;
Conservative (
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; encephalopathy; immune system stimulation.
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                                                                                                                                                                                 Spohn
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Pred. No. 1.1e-70;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                       07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00059902.
21-JAN-2002; 2002WO-IB00166.
19-JUL-2002; 2002US-0396636P.
                                                                                                                                                                                                                                                                                                                       antiallergic; antiasthmatic; cytostatic; vaccine; virus-like interleukin; IL-5; IL-13; eotaxin; repetitive antigen array; allergic eosinophilic disease; asthma; Hodgkin's lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a composition comprising a core particle having at least one first attachment site, and at least one antigen or antigenic determinant having at least one second attachment site. The antigen or antigenic determinant is a RANKL protein, RANKL fragment or RANKL peptide. The second attachment site is (non-) naturally occurring with the antigen or antigenic determinant, and is capable of association to the first attachment site. The antigen or antigenic determinant and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful as a medicament, or for the manufacture of a medicament for treating bone diseases. The composition is especially useful for as a vaccine for therapy or prophylaxis of bone diseases, particularly mammalian encephalopathies,
                                                                                 Bachmann
                                                                                                                                                                                                             07-NOV-2002; 2002WO-EP012455.
                                                                                                                                                                                                                                                                                                Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                 Virus-like particle
                                                                                                                                                                                                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                            ADK17144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADK17144 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 132 AA;
                                                                                                                                                                                                                                       15-MAY-2003.
                                                                                                                                                                                                                                                                     WO2003040164-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                             (CYTO-)
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                                                                                                             CYTOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    stimulating mammalian immune system. This sequence represents a of the invention.
                                                                                 Σ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LIDAIDQLNPAY 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AKLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LIDAIDQLNPAY 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AKLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                                                               Jennings
                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                             BIOTECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide; 132
                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                 repetitive antigen array peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                 Ģ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23; 222pp; English
                                                                               Sonderegger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 664; DB 7;
Pred. No. 1.1e-70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                    #23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                      particle;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       60
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Composition comprising an ordered and repetitive antigen or antigenic determinant array, useful as a medicament, or for manufacturing a medicament for treating allergic eosinophilic diseases, e.g. asthma, or

2003-441518/41.

antigen with at least one second attachment site, where the antigen is a protein or peptide of II-5, II-13 or entaxin. The second attachment site is an attachment site naturally or not naturally occurring with the antigen or antigenic determinant. The second attachment site is capable of association to the first attachment site, and where the antigen or antigenic determinant and the core particle interact through the association to form an ordered and repetitive antigen array. The association are useful as medicaments, or for manufacturing a medicament or a vaccine for treating allergic eosinophilic diseases, e.g. asthma, or Hodgkin's lymphoma and related diseases. This sequence is used to generate the compound of the invention.

The invention relates to a composition comprising a virus-like particle and at least one antigen, which is a protein or peptide of interleukin (IL)-5, IL-13 or ectaxin and is bound to the virus-like particle, or a core particle with at least one first attachment site and at least one

Disclosure; SEQ ID NO 23; 245pp; English

Hodgkin's lymphoma

S

1 AKLETYTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK

Query Match
Best Local Simi
Matches 132;

Similarity

100.0%; Score 664; DB 7; 100.0%; Pred. No. 1.1e-70;

Length 132;

0,

Conservative

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Mismatches

Sequence 132 AA;

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> 121 61 61

LIDAIDQLNPAY 132 LIDAIDQLNPAY 132

NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL AKLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK

120 120 60 60

RESULT 9

ADJ36313 standard; protein;

132

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New compositions comprising a virus-like particle (VLP), an immunostimulatory substance bound to the VLP, and an antiger the VLP, useful for enhancing immune response or for treatir
                                                                                                                                                                                                                                                          immune response; virus-li
chronic disease; chronic
                                                                                                                                                                                                                                                                                  antiallergic; cytostatic; virucide; immunostimulant; vaccine;
                                                                                                                                                                                                                                                                                                        Bacteriophage AP205 coat protein virus-like particle.
                                                                                                                                                                                                                                                                                                                                    22-APR-2004
                                                                                                                                               20-JUN-2003; 2003WO-EP006541.
                                                                                                                                                                         31-DEC-2003
                                                                                                                                                                                               WO2004000351-A1
                                                                                                                                                                                                                        Bacteriophage AP205
                                                  WPI; 2004-108361/11.
                                                                                                (CYTO-) CYTOS BIOTECHNOLOGY AG
                                                                                                                       20-JUN-2002; 2002US-0389898P.
                                                                         Renner
                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                        virus-like
                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                           ike particle; immunostimulatory; allergy; tumour;
viral disease; RNA bacteriophage; coat protein;
 and an antigen mixed or for treating e.g.
               with
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least one

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ADJ67159
III ADJ7
XX
AC ADJ7
XX
AC ADJ7
AC ADJ7
XX
Phag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 132; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 132 AA;
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The invention relates to a new composition comprising: (i) a core particle with at least one first attachment site; and (ii) at least antigen or antigenic determinant with at least one second attachment.
                                                                                                                                                                   New composition comprising a core particle having a first attachment and an antigen or antigenic determinant which is a ghrelin or ghrelin peptide having a second attachment site, useful for treating obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       anorectic; core particle;
antigenic array.
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                                                                                                                                                                                                                                                                                                                                                         Bachmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-JUL-2002; 2002US-0396638P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-JUL-2003; 2003WO-EP007849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteriophage Obeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Phage Qbeta coat protein mutant Qbeta 240 for antigen display array.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ67159;
                                                                                                                   Disclosure; SEQ ID NO 17; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LIDAIDQLNPAY 132
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                                                                                                                                                                                                                                                                                                                                                      MF, Fulurija
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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Pred. No. 1.1e-70;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            determinant; ghrelin; P-pilin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 132;
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                                                                                                                                                                               or ghrelin
g obesity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    site, where the antigen or antigenic determinant is ghrelin or a ghrelin peptide, and where the second attachment site being consisting of an attachment site not naturally occurring with the antigen or antigenic determinant and an attachment site naturally occurring with the antigen or antigenic determinant, where the second attachment site is capable of association to the first attachment site, and where the ghrelin or a ghrelin peptide and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful for treating obesity. The repetitive array may form part of a phage or bacterial display array. This peptide corresponds to a Bacteriophage obeta coat protein mutant which can used as part of the repetitive or
Composition comprising a core particle with at least one attachment site, and an antigenic amyloid beta 1-6 peptide, useful for treating diseases such as Alzheimer's disease.
                                                                                                                                               (CYTO-)
                                                                                                                                                                                               19-JUL-2002; 2002US-0396639P.
15-MAY-2003; 2003US-0470432P.
                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteriophage Obeta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           neuroprotective; nootropic; vaccine; amyloid beta 1-6 peptide; core particle; antigen array; Alzheimer's disease; RNA bacteriophage; coat protein; CP; mutant; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteriophage Qbeta coat protein mutant K13R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADK52193 standard;
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                                                                 WPI; 2004-203731/19.
                                                                                                               Bachmann MF,
                                                                                                                                                                                                                                              18-JUL-2003; 2003WO-EP007864.
                                                                                                                                                                                                                                                                                                                WO2004016282-A1
                                                                                                                                                                                                                                                                                                                                                               Misc-difference 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antigenic array.
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                                                                                                                                               CYTOS BIOTECHNOLOGY AG NOVARTIS PHARMA AG.
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                                                                                                             Tissot A,
                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                           /note= "Wild type Lys substituted by Arg"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 132
                                                                                                                 Ortmann
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                                                                                                               Staufenbiel
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RESULT 12
ABG94318
ID ABG94
XX ABG94
XX 29-AU
DT 29-AU
DT 10-DE
XX Cytos
XW Human
XW cytos
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC The invention describes a novel composition comprising a virus-like core CC particle with at least one attachment site, and an antigenic amyloid beta C1-6 peptide. The new composition comprises: a core particle with at least C1-6 cone first attachment site; and at least one antigen or antigenic C1-6 cone first attachment site; and at least one antigen or antigenic C1-6 cone first attachment site, and at least one antigen or antigenic C1-6 contains the site comprises an attachment site, where the antigen C1-6 countring with the antigenic determinant; or an attachment C1-7 cocurring with the antigenic determinant; or an attachment C1-7 cocurring with the antigenic determinant; or an attachment C1-7 contains the site is capable of association to the first attachment C1-7 composition is useful for the manufacture of a medicament for treating C1-7 composition is useful for the manufacture of a medicament for treating C1-7 composition is useful for the manufacture of a medicament for treating C1-7 composition of the compositions and vaccines of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 132; Conserv
                                   Renner WA,
Piossek C;
                                                                                                                                                                                                                                                                                                                                                              Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory; cytostatic; antiviral; antidiabetic; hypoglycaemic; antigen array; vaccine; infectious disease.
 WPI; 2002-627351/67.
                                                                                                                          19-JAN-2001; 2001US-0262379P.
04-MAY-2001; 2001US-0288549P.
05-CCT-2001; 2001US-032698P.
07-NOV-2001; 2001US-0331045P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 132 AA;
                                                                                                                                                                                                                                                                                                                              unidentified bacteriophage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-AUG-2003
10-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; SEQ ID NO 17; 184pp; English
                                                                                                                                                                                                                    21-JAN-2002; 2002WO-IB000166.
                                                                                                                                                                                                                                                        25-JUL-2002.
                                                                                                                                                                                                                                                                                            WO200256905-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                      pQB250 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABG94318 standard;
                                                                                        (CYTO-) CYTOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDBERAFVRTELAALLASPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AKLETVTLGNI GRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AKLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                      Bachmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(first entry)
                                                                                        BIOTECHNOLOGY AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 664; DB 8;
100.0%; Pred. No. 1.1e-70;
tive 0; Mismatches 0;
                                                    Σ
                                                    Tissot A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B
                                                      Maurer P,
                                                      Lechner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 132;
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ABG80630
ID ABG80
XX ABG80
AC ABG80
XX ABG80
XX Bacte
XX Molec
XX 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC the core particle by at least one covalent bond. Also disclosed is an CC antigen or antigenic determinant with at least one second attachment cC site, where the antigen or antigenic determinant is amyloid beta peptide (Abetal-42) or its fragment and where the second attachment site is selected from an attachment site not naturally occurring with the antigen cC antigenic determinant and an attachment site naturally occurring with the antigen or antigenic determinant, where the second attachment site is cC capable of association through at least one non-peptide bond to the first cC attachment site and where the association to form an ordered and cC repetitive antigen array. The invention also comprises a coat protein cC capable of forming a capsid which comprises mutant Obeta coat protein cC capable of forming a capsid which comprises mutant Obeta coat protein cC capable of forming a capsid which comprises mutant Obeta coat protein cC capable of forming a capsid which comprises mutant Obeta coat protein cC capable of forming a capsid which comprises mutant Obeta coat protein cC capable of forming a capsid which compounds of the invention may compound to the specification. The compounds of the invention may construct the second attachment and the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                 Molecular antigen array; vaccine; antigen; antimicrobial; molecular scaffold; amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgB-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus erythematosus; osteoporosis; influence disease; myasthenia gravis; multiple sclerosis; immunoproliferative disease; myasthenia gravis; multiple sclerosis; angioimmunoproliferative disease; lymphadenopathy; Alzheimer's disease; angioimmunoproliferative lymphadenopathy; immunoblastive lymphadenopathy; rheumatoid arthritis; diabetes; infectious disease; factor Xa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprising a core particle selected from a core particle of a non-natural origin and a core particle of natural origin and an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond. Also disclosed is an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nave antimicrobial, antiallergic, immunomodulatory, cytostatic, antiviral, antidiabetic, or hypoglycaemic activities and may be used in immunisation and as a vaccine. The present sequence represents a proteir sequence used to create the compositions of the invention. (Updated on 2-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bacteriophage Q-beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            also
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecular antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABG80630 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention relates to a novel ordered and repetitive antigen array in the production of vaccines for infectious diseases. The invention discloses a composition comprising a non-natural malacritical material mate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                  cysteine-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     array used in the production of vaccines for infectious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Al coat protein mutant Qbeta-250.
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1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 132 AA
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                                                                       linker
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.4e-70;
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CC one second attachment site, where the antigen or antigenic determinant is attachment site is selected from: (i) an attachment site not naturally occurring with the antigen or antigenic determinant is occurring with the antigen or antigenic determinant, and (ii) an attachment site not naturally occurring with the antigen or antigenic determinant, and (ii) an occurring with the antigen or antigenic determinant, and (ii) and continuous attachment site is capable of association occurring with the antigen or antigenic determinant and the scaffold interact through at least one non-peptide bond to the first attachment site; and through at least one non-peptide bond to the first attachment site; and continuous attachment site is capable of association occurring through the association to form an ordered and repetitive antigen array. The composition is used in continuous attachment site antigen array. The composition is used in continuous disease, influence, graft or producing a non-naturally occurring continuous disease, influence, graft or producing a non-naturally occurring continuous disease, influence, graft or producing a non-naturally occurring continuous disease, influence, graft or producing a non-naturally occurring continuous disease, influence, graft or producing a non-naturally occurring continuous disease, influence, graft or producing a non-naturally occurring continuous disease, influence, graft or producing a non-naturally occurring continuous diseases, graft or producing a non-natu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          molecular scaffold comprising: (i) a core particle selected from: (1) a core particle of a non-natural origin; and (2) a core particle of natural origin; and (12) a core particle of natural origin; and (14) an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond; (b) an antigen or antigenic determinant with at least one covalent bond; (b) an antigen or antigenic determinant with at least
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Maurer
Renner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteriophage Qbeta.
Synthetic.
                                          or C-terminal linker peptide which serves as the attachment virus like particle or bacterial protein (the scaffold prote present sequence is bacterial protein or peptide which is co modified antigen to form the molecular antigen array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JAN-2001; 2001US-0262379P.
04-MAY-2001; 2001US-0288549P.
05-CCT-2001; 2001US-032698P.
07-NOV-2001; 2001US-0331045P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a composition comprising: (a) a non molecular scaffold comprising: (i) a core particle selected
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              The present invention describes a composition (C) for enhancing an immune CC response against an antigen or a virus-like particle in an animal. (C) CC comprises a virus-like particle (VIP) bound to at least one antigen, or a CC VIP capable of being recognised by the immune system of the animal. Also CC described: (1) enhancing an immune response against an antigen or a VIP CC in an animal comprising introducing (C) into the animal; (2) vaccines CC comprising (C) together with a pharmaceutical diluent, carrier or CC exciptent; (3) immunising or treating an animal comprising administering CC the vaccine to the animal, or priming or boosting a T cell response in CC the animal by administering the vaccine; and (4) enhancing anti-viral CC protection in an animal comprising introducing (C) into the animal. (C) CC has cytostatic, virucide, antibacterial, antiparasitic, fungicide, antitallergic, immunosuppressive, antiaddictive, antiaflammatory, CC antitheumatic and antiarthritic activities. (C) or the vaccines can be used for enhancing antiaviral parotection in an animal, or immunising or canimal, enhancing antiaviral protection in an animal, or immunising or canimal, enhancing antiaviral protection in an animal, or immunising or canimal, antipararial and informations and information and control are viral bacterial.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A composition, useful for enhancing or a virus-like particle, enhancing or immunizing or treating tumors or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 181; 243pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-SEP-2002; 2002WO-IB004252
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treating
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diseases such as
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anti-viral protection in an animal,
infectious diseases, e.g. viral
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-IB000166.
19-JUL-2002; 2002US-0396637P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteriophage Qbeta.
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The invention describes an angiotensin peptide moiety carrier conjugate comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second

Claim 20; Page 94; 97pp; English

New angiotensin peptide moiety carrier conjugate comprising a carrier and an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension

និនិនិនិនិនិនិនិង Ś 밁 Ş 밁 문 Query Match 99.5%; Score 661; DB 6; Length 132; Best Local Similarity 99.2%; Pred. No. 2.4e-70; Matches 131; Conservative 1; Mismatches 0; Indels attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kidney failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta coat protein used in the preparation of the vaccine conjugates of the invention Sequence 132 AA; 121 LIDAIDQLNPAY 132 ||||||||| 121 LIDAIDQLNPAY 132 61 NYKVQVKIQNFTACTANGSCDFSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120 61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120 0, Gaps 0

Search completed: January 4, 2005, 09:11:00 Job time: 40.6824 secs

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4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
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US-08-09-148-185
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RESULT 2
US-09-948-722-2
US-09-948-722-2;
Sequence 2, Application US/099487;
Patent No. 6776993;
GENERAL INFORMATION:
APPLICANT: Kaufmann, Stefan H. E.
APPLICANT: Hess, Jurgen
; TITLE OF INVENTION: Tuberculosis

US/09948722

Tuberculosis Vaccine

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RDGKOTLVLNP	LENGTH: 626 TYPE: PRI ORGANISM: Artificial FEATURE: OTHER INFORMATION: D 9-485-717-2 9-485-717-2 ery Match st Local Similarity tches 29; Conserv	APPLICANT: Kaufmann, Stefan APPLICANT: Hess, Jrgen Tifle OF INVENTION: Tubescu FILE REFERENCE: 16862PUS CURRENT APPLICATION NUMBER: ENGINE APPLICATION NUMBER: ENGINE FILING DATE: 1997-08-PRIOR APPLICATION NUMBER: ENGINE FILING DATE: 1998-08-PRIOR APPLICATION NUMBER: ENGINE FILING DATE: 1998-08-PRIOR FILING DATE: 1998	7-2 Applicati 6673353 FORMATION:	9.9.9.9
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ELVENOPDVLPVKRDSL 197QKYADVTFSFTQYSTD 102	nce: recombinant Length 626; Indels 35; Gaps 5;			Sequence 2885, Apple Sequence 76, Apple Sequence 775, Apple Sequence 41, Apple Sequence 13, Apple Sequence 29602, A Sequence 25, Apple Sequence 35, Apple Sequence 37, Apple Sequence 26, Apple Sequence 27, Apple Sequence 28, Apple Sequence 27, Apple Sequence 28, Apple Sequence 29, Apple Sequence 21, Apple Sequence 12, Apple Sequence 12, Apple Sequence 5, Appli

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PRIOR FILING DATE: 1998-08-12
PRIOR APPLICATION NUMBER: EP 97/114,614.7
PRIOR FILING DATE: 1997-08-23
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
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GENERAL INFORMATION:
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SE
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND
                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: GTC99-03PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/485,717
PRIOR FILING DATE: 2000-02-22
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                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Acinetobacter baumannii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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ORGANISM: Artificial Sequence
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                                                                                                                                                                           13 RDGKQTLVLNPR-----GVNPTNGVASLSQAGAVPALEKRVTV-SVSQPSRNRKNYKVQ 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13 RDGKQTLVL-----NPRGVNPTNGVASLSQAGA---
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                                                                                                                                                                                                                                                                                                                                           409
                                                                                                                                                                                                                                  Similarity
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                                   DQL 128
                                                                                                       VKIONPTACTANGSCDPSVTROKYADVTFSFTQYSTDEERAFVRTELAALLASPLLIDAI 125
                                                                                                                                             RGGTRLNVNVPRNALISFAVNPVDGTTPI----IAKPNQRLTIRDLLAPGRTGSN---A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EERAFVRTELAA 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TVSVSQPSRNRKNYKVQVKIQNPTACTANGSCDPSVTR-----QKYADVTFSFTQYSTD 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09328352
                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                  Score 75; DB 4; Length 409; Pred. No. 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 81.5; D
Pred. No. 0.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 0.3;
4; Mismatches
                                                                                                                                                                                                                   Mismatches
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                                                                       -EEVMESVKTIAHMLKASKQILDDL
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                                                                       222
                                                                                                                                             169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 27
                                                                                                                                                                                                                                                                                                     APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEG ID NOS: 62517
SOFTWARE: PATENTIN Ver. 2.0
SEG ID NO 44620
LENGTH: 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 23637, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 23637
LENGTH: 916
                                                                                                                                                Matches
                                                                                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 44620, Appearent No. 6703491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 1998-02-18 PRIOR APPLICATION NUMBER: US PRIOR PILING DATE: 1998-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATITIES OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE:
                                                                                                                                                                                                                                                          ORGANISM: Drosophila melanogaster FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
                                                                                                                                                                                                                                       OTHER INFORMATION: Xaa means any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       789 GLLGGEGALARLRAAIVVQARRIDPFQGLAAALPVGGLPALEQAEDFAFGGSLDQRWVQA 848
                                                                     849 EERGAEYRAQESAHREHPCRSIEANADCPLSADAGKW-----PWTIFSDSATRWALRT 901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
27; Conserv
                                                                                                          18 TLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQP-----SRNRKNYKVQVKIQN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    56 SRNRKNYKVQVKIQNPTAC---TANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRT 110
                                                                                                                                            29;
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                                                                                                                                                                 Similarity
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                                  PTACTANGSCDPSVTRQKYADVTFSFTQ-----YSTDEERAFVRT--ELAALLASPLL 121
AXXVVKYFSCTSSATRHSTRCISSYVSQSLGPSPLLGXADXKDFVRTLXPLATFISSPNL
                                                                                                                                                Conservative
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NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
                                                                                                                                                                 10.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33142
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22.9%; Pred. No. 10;
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                                                                                                                                            20;
                                                                                                                                                                 Score 70.5;
Pred. No. 5.
                                                                                                                                                Mismatches
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RESULT 6 US-08-090-148-1

Sequence 1, Application US/08090148 Patent No. 5534257

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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 940-8636
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 AMINO ACIDS
                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/
FILING DATE: 08/11/93
CCLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 91
FILING DATE: 01/24/91
APPLICATION NUMBER: PCT/GB
FILING DATE: 01/22/92
ATTORNEY/AGENT INFORMATION:
NAME: Nissenbaum, Israel
REGISTRATION NUMBER: 27,58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: NOT MOLECULE TYPE: PORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                   PUBLICATION: SEQUENCE:
PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 10022-2585
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: S. York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NITLE OF INVENTION: Antigen-Presenting Capsid with ITLE OF INVENTION: Fusion MS2-Coat Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PS2-486
OPERATING SYSTEM: MS-
SOFTWARE: WordPerfect
                                                                                                                                                                                                                   PAGES: 82-88
DATE: 1972
                                                                                                                                                                                                                                                                JOURNAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
77 LPVAAWRSYLNMELTIPIFATNSDCELIVKAMQGLL 112
                               81 DPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALL 116
                                                                24 NFANGVAEWISSNSRSQA-----YKVTCSVRQSSAQNRKYTIKVEVPKVATQTVGGVE 76
                                                                                               27 NPTNGVA-----SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVKIQNPTACTANGSC 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : 129 AMINO ACIDS
AMINO ACID
GY: NOT RELEVANT
                                                                                                                                                                                                                                                                                  bacteriophage MS2 coat protein
                                                                                                                                                                                                                                                                                                 : Ysebaert, M.; Piers, W. Nucleotide sequence of the gene coding for the
                                                                                                                                                                                                                                                                                                                                  Min Jou, W.; Haegeman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                575 Madison Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mastico, Robert Allan
Stockley, Peter George
Talbot, Simon John
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                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                  Nature
                                                                                                                                                                                                                                                                                                                                                                                             Location 1 through 129 below represents entire MS2 coat protein in the published
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Coat protein from MS2-RNA
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                                                                                                                                                10.5%;
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                                                                                                                                  16;
                                                                                                                                Score 69.5; DB Pred. No. 0.85; 6; Mismatches
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                                                                                                                                                               DB 1;
                                                                                                                                  43; Indels
                                                                                                                                                               Length 129;
                                                                                                                                13;
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Query Match
Best Local Similarity
Matches 38; Conserve
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US-09-302-626B-182
                                                                                         US-09-302-626B-185
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PRIOR FILING DATE: 1999-01-14
NUMBER OF SEQ ID NOS: 195
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 182
                                                                                                                                                                                               PILE REPERENCE: CHIROLS9
CURRENT PILLING DATE: 199-04-30
PRIOR APPLICATION NUMBER: PCT/IB99/00103
PRIOR FILING DATE: 1999-01-14
NUMBER OF SEQ ID NOS: 195
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATI
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Pizza, Mariagrazia
APPLICANT: Grandi, Guido
TITLE OF INVENTION: Meningococcal Antigens
FILE REFERENCE: CUIRO159
CURRENT APPLICATION NUMBER: US/09/302,626B
CURRENT FILING DATE: 1999-04-30
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Pizza, Mariagrazia
APPLICANT: Grandi, Guido
TITLE OF INVENTION: Meningococcal Antigens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-302-626B-185
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APPLICANT: Masignani, Veg
APPLICANT: Rappuoli, Rino
                                                                                                                                                                  LENGTH: 62
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Scalato, Enzo
APPLICANT: Masignani, V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                           OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                            FEATURE:
                                                                                                                                          ORGANISM: Artificial Sequence
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o. 6709660
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5. 6709660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     50 VSVSQPSRNR-----KNYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDE 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10 NIGRDGKQTLVLNPRGVNPTNGVASLSQAGAV------
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Rappuoli, Rino
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               10.4%; Score 69; DB ilarity 23.2%; Pred. No. 12; Conservative 17; Mismatches
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23.2%; Pred. No.
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                                     DB 4;
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                                                Length 622;
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                   Gaps
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Polypeptide Accession Number YDR490C
US-09-538-092-216
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US-09-206-942-49
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PRIOR FILING DATE: 2000-02-01
NUMBER OF SEQ ID NOS: 1387
SOFTWARE: CuraPatSeqFormatter Version 0.9
SEQ ID NO 216
LENGTH: 766
TYPE: PRT
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GENERAL INFORMATION:
                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Loosmore, Sheena M.
                                                                                                                                                                                                                                                Sequence 49, Application US/09206942
Patent No. 6432669
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APPLICANT: Mansfield, Traci A.
APPLICANT: Mansfield, Traci A.
TITLE OF INVENTION: Protein-Protein Complexes
FILE REFERENCE: 15966-542
CURRENT APPLICATION NUMBER: US/09/538,092
CURRENT FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: 60/127,352
PRIOR FILING DATE: 1999-04-01
APPLICANT: Yang, Yan-Ping
APPLICANT: Klein, Michel H.
TITLE OF INVENTION: Protective Recombinant Haemophilus Influenzae High
TITLE OF INVENTION: Molecular Weight Proteins
FILE REFERENCE: 1038-861 MIS:jb
CURRENT APPLICATION NUMBER: US/09/206,942
CURRENT FILING DATE: 1998-12-08
EARLIER APPLICATION NUMBER: 09/167,568
EARLIER FILING DATE: 1998-10-07
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RESULT 12
US-09-198-452A-745
(Sequence 745, Application US/09198452A)
Patent No. 6559294
GENERAL INFORMATION:
APPLICANT: Griffais, R.
TITLE OF INVENTION: thereof and uses the control of 
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; TYPE: PRT
; ORGANISM: Haemophilus influenzae
US-09-206-942-47
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 49
LENGTH: 1073
TYPE: PRT
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EARLIER APPLICATION NUMBER: 09/167,568
EARLIER FILING DATE: 1998-10-07
NUMBER OF SEQ ID NOS: 95
SOFTWARE: Patentin Ver. 2.1
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CURRENT FILING DATE: 1998-12-08
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TITLE OF INVENTION: Molecular Weight Proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         55 PSRN---RKNYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTE
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27; Conserv
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27; Conserv
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22.0%;
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22.0%; Pred. No.
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Pred. No.
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a pneumoniae genomic sequence and polypeptides, and uses thereof, in particular for the diagno

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Best Local Similarity
Thes 26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; NAME/KEY: SITE
; LOCATION: 1...381
; OTHER INFORMATION:
US-09-198-452A-745
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-045-186-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/09045186 Patent No. 6087154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 745
LENGTH: 381
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: 1
CURRENT FILING DATE: 1998-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                         TELEFAX: (317) 276-3861 (NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                    TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 383 amino acids
                                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Gavic Band I
                                                                                                                                                                         NAME: Gaylo, Paul J.
REGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: P-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 276.0756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Baez, Melvyn
APPLICANT: Cates, Carolyn A.
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Liii
                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VKTVIL-NIGRKILQGIKKXKKKIGIXSGLFFLDLV-----LLGVSXQRPTETSANV 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KVQVKIQNPTACTANGSCDPSVTRQKYADVTFS----FTQYSTDEBRAFVRTELAALL 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indiana
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Lill Corporate Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  United States of America
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                                                                        linear
                                                    protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10.3%; Score 68.5; DI
19.8%; Pred. No. 6.3;
10.3%;
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Score 68.5;
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Length 383;
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Patent No. 5
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                                                                                                                                                                                                                            Query Match
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APPLICANT: SELBIE
                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 202-783-6040
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: HERZOG, Herbert
APPLICANT: SHINE, John
TITLE OF INVENTION: Human Neuropeptide Y-Y1 Receptor
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: ERNST, Barbara G
REGISTRATION NUMBER: 30,377
                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 26 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
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                                                                                                                                                                                                                                                                                                                                                      ENGTH:
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                                                                                                                      138 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN-
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                                                                                   68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
                                                                                                                                                         16 KQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
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555 13th St, N.W., Suite 701-East
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SELBIE, Lisa
HERZOG, Herbert
SHINE, John
                                                                                                                                                                                                                                                                                                                                                    384 amino acids
                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                protein
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                                                                                                                                                                                                           10.3%; Score 68.5; DB 1; 25.0%; Pred. No. 6.4;
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                                                   -VTLDAYKDKYVCFDQFPSDSHRLSYTTLLLVL 218
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; Patent No. 5958709 ; GENERAL INFORMATION: ; APPLICANT: Bard, J ; APPLICANT: Walker,

Bard, Jonathan A Walker, Mary US-08-555-268A-15

Sequence 15, Application US/08555268A

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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC COMPOTER:
COMPUTER: COMPOTER: PATENTH POLOS/MS-DOS
SOFTWARE: PATENTH Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/555,268A
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 4743-Z/JPW/MAT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-04400
TELEPAX: (212) 391-0526
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 384 amino acids
TYPE: amino acid
STRANDEDNESS: single
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
Search completed: January 4, 2005, 09:35:43
Job time: 11.3622 secs
                                                                                                                                                                                                                                                                                                               Query Match 10.3%; Score 68.5; DB 2; Length 384; Best Local Similarity 25.0%; Pred. No. 6.4; Matches 27; Conservative 12; Mismatches 34; Indels 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Branchek, Theresa
APPLICANT: Weinshank, Richard L.
TITLE OF INVENTION: DNA ENCODING A HUMAN NEUROPEPTIDE
TITLE OF INVENTION: Y/PEPTIDE Y/PANCREATIC POLYPEPTIDE
TITLE OF INVENTION: RECEPTOR (Y4) AND USBS THEREOF
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 1185 Aver CITY: New York STATE: New York COUNTRY: U.S.A. ZIP: 10036
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                                                                                                                                                      68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTPEERAFVRTELAAL 115
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1185 Avenue of the Americas
                                                                                                         -VTLDAYKDKYVCFDOFPSDSHRLSYTTLLLVL 218
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Searched:
                                                                                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB DB
                                                                                                                                                                                                                                                  Score
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Match
     100.0
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Gapop 10.0 , Gapext 0.5
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664
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112:
113:
114:
116:
116:
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                                                                                                                                                                                                                                                                                                                                                                                                 : /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
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: /cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-10-243-739-23

US-10-244-055-23

US-10-289-454-23

US-10-050-902-255

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US-10-346-190-23

US-10-346-191-14

US-10-465-811-14

US-10-622-064-6

US-10-622-064-7

US-10-622-124-17

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US-10-622-124-17

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US-10-622-057-17

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1450.557 Million cell updates/sec
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                                    23, Appl
23, Appl
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255, Appl
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IPAY 132	NPTACT/	NIGRDGI NIGRDGI	100. Y 100. rvative	ication US/10243 S20030091593A1 ON: ann, Martin F. ni, Tazio ni, Tazio con: Franziska ON: In vivo Acti ON: Immune Resp 1700.0210001 ION NUMBER: US/1 ATE: 2002-09-16 N NUMBER: 60/318 E: 2001-09-14 NOS: 73 In version 3.1 riophage Q-beta	1132 1132 1132 1132 1132 1132 1132 1132
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	NYKVQVKIQNPTACTANGSCDPSVTRQKVADVTFS 	VLNPRGVNPTN VLNPRGVNPTN	Score 664 Pred. No. 0; Mismatc	ALIGN 739 vation onses 0/243,	US-10-289-454-25 US-10-050-902-257 US-10-346-190-25 US-10-465-811-16 US-10-289-456-25 US-10-622-064-9 US-10-622-124-19 US-10-622-124-19 US-10-622-087-19 US-10-244-065-26 US-10-10-249-454-26 US-10-10-465-811-17 US-10-622-064-9 US-10-243-739-10 US-10-244-065-24 US-10-244-065-27 US-10-244-065-29 US-10-249-454-24 US-10-299-454-24 US-10-299-454-24 US-10-050-902-256 US-10-050-902-256
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	FTQYSTDEERAFVRTELAALLASPL 120 	AKLETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAĠAVPALEKRVTVSVSQPSRNRK 60 	Length 132; Indels 0; Gaps 0;	Presenting Cells for Enhancement of Virus Like Particles	Sequence 25, Appl Sequence 257, App Sequence 257, App Sequence 25, Appl Sequence 16, Appl Sequence 18, Appl Sequence 18, Appl Sequence 19, Appl Sequence 26, Appl Sequence 27, Appl Sequence 26, Appl Sequence 27, Appl Sequence 27, Appl Sequence 27, Appl Sequence 27, Appl Sequence 20, Appl Sequence 20, Appl Sequence 20, Appl Sequence 21, Appl Sequence 21, Appl Sequence 22, Appl Sequence 24, Appl Sequence 27, Appl Sequence 28, Appl Sequence 28, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 27, Appl Sequence 28, Appl Sequence 28, Appl Sequence 27, Appl Sequence 27, Appl Sequence 28, Appl Sequence 27,

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US-10-244-065-23
                                                                                                                                                                                                                                                                     Sequence 23, Application US/10289454
Publication No. US20030157479A1
GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
LENGTH: 132
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APPLICANT: Stormi, T
APPLICANT: Maurer, P
APPLICANT: Tissot, A
APPLICANT: Schwarz, I
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              TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
CURRENT APPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/1802/00166
PRIOR APPLICATION NUMBER: PCT/1802/00166
PRIOR FILING DATE: 2002-01-21
PRIOR PILING DATE: 2002-01-18
PRIOR FILING DATE: 2002-01-18
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APPLICANT: Jennings, Gary
APPLICANT: Sonderegger, Ivo
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APPLICANT:
APPLICANT:
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APPLICATION NUMBER: US 60/331,045
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132; Conserv
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||IDAIDQLNPAY 132
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Cielens, Indulis
Renhofa, Regina
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Sonderegger, I
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Lipowsky, Gerad
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  LIDAIDQLNPAY 132
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; LENGTH: 132
TYPE: PRT
; ORGANISM: Bacteriophage Q-beta-240
US-10-289-454-23
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 23
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SEQ ID NO 255
LENGTH: 132
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Best Local Similarity
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Best Local Similarity
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CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR FILING DATE: 2001-01-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lechner, Franziska
APPLICANT: Sebbel, Peter
APPLICANT: Piossek, Christine
TITLE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0190004
                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/331,045 PRIOR FILING DATE: 2001-11-07
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PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
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121 LIDAIDQLNPAY 132
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                                           61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                            61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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to. US20030175290A1
                                                                                                                                                                                                                    100.0%; Score 664; DB 14; ilarity 100.0%; Pred. No. 2.1e-66; Conservative 0; Mismatches 0:
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US-10-050-898-255

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APPLICANT: Bachmann, Martin
APPLICANT: Maurer, Patrick
APPLICANT: Maurer, Patrick
APPLICANT: Pellicioli, Erica
APPLICANT: Renner, Wolfgang A.
TITLE OF INVENTION: Prion Protein Carrier-Conjugates
FILE REFERENCE: 1700.029003
CURRENT APPLICATION NUMBER: US/10/346,190
CURRENT FILING DATE: 2003-01-17
PRIOR APPLICATION NUMBER: 60/396,590
PRIOR PILING DATE: 2002-07-18
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/389,898
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US-10-346-190-23
US-10-346-190-23
Sequence 23, Application US/10346190
Publication No. US20030219459A1
GENERAL INFORMATION:
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US-10-050-898-255
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PRIOR FILLING DATE: 2001-01-19
PRIOR PELLING DATE: 2001-05-04
PRIOR PILLING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILLING DATE: 2001-10-05
PRIOR PILLING DATE: 2001-11-07
PRIOR FILLING DATE: 2001-11-07
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SOFTWARE: PatentIn Ver
SEQ ID NO 255
LENGTH: 132
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Ortmann, Rainer
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Sebbel, Peter
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Maurer, Patrick
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100.0%; Pred. No. 2.1e-66;
vative 0; Mismatches 0;
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; ORGANISM: Qb
US-10-346-190-23
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                                                                                                                                                                                                                        Matches 132; Conservative
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Best Local Similarity
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Best Local Similarity
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CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: US 60/389,898
PRIOR FILING DATE: 2002-06-20
NUMBER OF SEQ ID NOS: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BACHMANN, MARTIN F
APPLICANT: RENNER, WOLFGAME A
TITLE OF INVENTION: PACKAGED VIRUS-LIKE PARTICLES FOR USE AS ADJUVANTS:
TITLE OF INVENTION: METHOD OF PREPARATION AND USE
FILE REFERENCE: 1700.0290004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: 10/050,902
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NUMBER OF SEQ ID NOS: 164
                                                                                                                                                                                                                                                                                                                                      LENGTH: 132
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Bacteriophage Qbeta 240 mutant
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                             121 LIDAIDQLNPAY 132
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                                                                      61 NYKVOVKIONPTACTANGSCDPSVTROKYADVTFSFTQYSTDEBRAFVRTELAALLASPL
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LIDAIDQLNPAY 132
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Pred. No. 2.1e-66;
D; Mismatches 0;
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US-10-289-456-23 ; Sequence 23, Application US/10289456

Publication No. US20040033211A1 GENERAL INFORMATION:

APPLICANT: Bachmann, Martin

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                                                                      ; OTHER INFORMATION: Bacteriophage Qbeta 240 mutant US-10-622-064-6
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Publication No. US20040059094A1
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Best Local Similarity
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Best Local Similarity Matches 132; Conserv
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PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 170
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PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR FILING DATE: 2002-01-18
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CURRENT FILING DATE: 2002-11-07
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APPLICANT: Spohn, Gunther
TITLE OF INVENTION: Antigen Arrays for Treatment of Bone Disease
FILE REFERENCE: 1700.0330001
                   Query Match
Best Local :
                                                                                                                                                                                 SEQ ID NO 6
                                                                                                                                                                                                                                                                      APPLICANT: Bachmann, Martin F
APPLICANT: Maurer, Patrik F
TITLE OF INVENTION: Hapten-Carrier Conjugates and Uses Thereof
FILE REFERENCE: 1700.0300001
CURRENT APPLICATION NUMBER: US/10/622,064
CURRENT FILING DATE: 2003-07-18
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PRIOR FILING DATE: 2002-07-18
NUMBER OF SEQ ID NOS: 33
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.2
                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                         FEATURE:
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                 100.0%; Score 664; DB 15; 100.0%; Pred. No. 2.1e-66;
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Pred. No. 2.1e-66;
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     Mismatches
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US-10-622-087-17
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,638
PRIOR FILING DATE: 2002-07-19
NUMBER OF SEQ ID NOS: 146
SOFTWARE: PatentIn version 3.2
SEQ ID NO 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17, Application US/10622124 Publication No. US20040076645A1 GENERAL INFORMATION:
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Best Local Similarity
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SOFTWARE: PatentIn version 3.2 SEQ ID NO 17
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APPLICANT: Fulurija, Alma
TITLE OF INVENTION: Ghrelin-Carrier Conjugates
FILE REFERENCE: 1700.0340001
                                   TITLE OF INVENTION: Amyloid Beta 1-6 Antigen Arrays
FILE REFERENCE: 1700.0350002
CURRENT APPLICATION NUMBER: US/10/622,087
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,639
PRIOR FILING DATE: 2002-07-19
PRIOR PPLICATION NUMBER: US 60/470,432
PRIOR APPLICATION NUMBER: US 60/470,432
PRIOR PILING DATE: 2003-05-15
PRIOR FILING DATE: 2003-05-15
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APPLICANT: Tissot, Alain
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TYPE: PRT
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                                                                                                                                                                                                                                 Frey, Peter
                                                                                                                                                                                                                                                Staufenbiel, Matthias
                                                                                                                                                                                                                                                                          Luond,
                                                                                                                                                                                                                                                                                              Ortmann, Rainer
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                                                                                                                                                                                                                                                                             Rainer
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Pred. No. 2.1e-66;
Mismatches 0;
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NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 25
LENGTH: 132
TYPE: PRI
OPENATOR:
                                                                                                                               RESULT 13
US-10-244-065-25
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US-10-243-739-25
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; ORGANISM: Bacteriophage Q-beta
US-10-243-739-25
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APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Lechmer, Franziska
TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
FILE REFERENCE: 1700.0210001
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
PRIOR FILING DATE: 2001-09-14
                                                                        Sequence 25, Applica Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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Best Local Similarity
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                 APPLICANT: Bachmann,
APPLICANT: Storni,
APPLICANT: Maurer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Bacteriophage Qbeta 240 mutant
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TYPE: PRT
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Storni, Tazio
Maurer, Patrick
Tissot, Alain
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                                                                                                           Application US/10244065
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                                                                                             US20030099668A1
                                                          Martin F
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99.2%; Pred. No. 4.5e-66;
tive 1; Mismatches 0; Indels
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APPLICANT: Schwarz, Katrin
APPLICANT: Meijerink, Edwin
APPLICANT: Meijerink, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Cielens, Indulis
APPLICANT: Cielens, Indulis
APPLICANT: Renhofa, Regina
TITLE OF INVENTION: Method of Preparation and Use
FILE REFERENCE: 1700.0220001
CURRENT APPLICATION NUMBER: US/10/244,065
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
NUMBER OF SEQ ID NOS: 73
                                                                                              ; TYPE: PRT ; ORGANISM: Bacteriophage Q-beta-250 US-10-289-454-25
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; ORGANISM: Bacteriophage Q-beta
US-10-244-065-25
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                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 386
SOPTWARE: PatentIn version 3.2
SEQ ID NO 25
LENGTH: 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25, Application US/10289454 Publication No. US20030157479A1 GENERAL INFORMATION:
  Query Match
Best Local Similarity
Matches 131; Conserv
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Best Local Similarity
                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR PILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/331,045
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bachmann, Martin
APPLICANT: Jennings, Gary
APPLICANT: Jennings, Gary
APPLICANT: Sonderegger, Ivo
APPLICANT: Sonderegger, Ivo
FITE OF INVENTION: Antigen Arrays for Treatments of Allergic Bosinophilic Diseases
FILE REFERENCE: 1700.0360001
CURRENT EPPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
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    Conservative
                        99.5%;
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Score 661; DB 14;
Pred. No. 4.5e-66;
1; Mismatches 0;
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APPLICANT: Maurer, Patrick
APPLICANT: Lechner, Franziska
APPLICANT: Sebbel, Peter
APPLICANT: Sebbel, Peter
APPLICANT: Piossek, Christine
FILE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0190004
CURRENT APPLICATION NUMBER: US 60/262,379
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-10-05
PRIOR FILING DATE: 2001-10-05
PRIOR FILING DATE: 2001-11-07
NUMBER: DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 350
SOFTMARE: Patentin Ver. 2.1
SEQ ID NO 257
LENGTH: 132
TYPE: PRT
ORGANISM: Ob 250
US-10-050-902-257
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US-10-050-902-257
; Sequence 257, App
; Publication No. U
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Search completed: January 4, 2005, 09:41:21 Job time: 32.7351 secs
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Matches 131; Conserv
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Tissot, Alain
Maurer, Patrick
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o. US20030175290A1
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Copyright

Run

Sequence:

Searched:

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Minimum DB
Maximum DB
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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          529.5
655.5
116.8
87.5
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81.5
81.5
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Result No.

Database

RESULT 2 \$01964 readthrough protein - phage SP C;Species: phage SP C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #	Qy 61 NYKVQVKIQNETACTANGSCDESVTRQKYADVTESETQYSTDEERAFVRTELAALLASPL	Query Match 98.6%; Score 655; DB 1; Length 132; Best Local Similarity 98.5%; Pred. No. 2.6e-57; Matches 130; Conservative 1; Mismatches 1; Indels 0; Gaps Matches 130; Conservative 1; Mismatches 1; Indels 0; Gaps Oy 1 AKLETYTIGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK	ewiD:557	A;Molecule type: mRNA A;Residues: 1-80 <esc> A;Cesidues: 1-80 <esc> A;Cross-references: UNIPROT:P03615 A;Croll, E.; Wilson, K.J.; Reiser, J.; Weissmann, C. J. Biol. Chem. 252, 990-993, 1977 J. Biol. Chem. 252, 990-993, 1977 A;Ticle: Revised amino acid sequence of Qbeta coat protein between positions A;Reference number: A92221; MUID:77118576; PMID:838709 A;Accession: A92221</esc></esc>	RESULT 1 VCBPQB coat protein - phage Q-beta C;Species: phage Q-beta C;Species: phage Q-beta C;Date: 29-Jul-1981 #sequence revision 24-Sep-1981 #: C;Accession: A92240; A92221; Ā92088; A04224 R;Escarmis, C.; Sastry, P.A.; Billeter, M.A. J; Biol. Chem. 253, 8390-8399, 1978 J; Biol. Chem. 253, 8390-8399, 1978 A;Title: Determination of the first half of the coat A;Reference number: A92240; MUID:79048469; PMID:3617- A;Accession: A92240	ALIGNMENTS	30 73.5 11.1 719 2 T39271 31 73 11.0 243 1 VHVUPT 32 73 11.0 430 2 G88884 33 72.5 10.9 129 1 VCBPR7 34 72.5 10.9 129 1 VCBPR7 35 72.5 10.9 129 1 VCBPR7 36 72.5 10.9 289 2 C70400 37 72 10.8 248 2 A86786 38 72 10.8 1461 2 E90696 38 72 10.8 1461 2 E90696 40 71.5 10.8 640 2 S48423 41 71 10.7 396 2 C84316 42 71 10.7 397 1 A43685 43 71 10.7 463 2 S00678 44 70.5 10.6 191 2 H90078 45 70 10.5 1545 2 T42751
text_change 09-Jul-2004	YSTDEERAFVRTELAALLASPL 120 	1; Length 132; 1; Indels 0; Gaps 0; 1; Indels 0; Gaps 0; LSQAGAVPALEKRVTVSVSQPSRNRK 60	protein.	rotein between positions 1 and 60.	#text_change 09-Jul-2004 t protein cistron of bacteriophage Qbe		conserved hypothet nucleocapsid prote protein K09B11.10 coat protein - pha coat protein - pha coat protein - pha coat protein - pha ferredoxin oxidore conserved hypothet hypothetical prote hypothetical prote hypothetical prote hypothetical prote polymerase-associa translation elonga hypothetical prote sulfonylurea recep

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R;bhaese, P.; Vandekerckhove, J.S.; Van Montagu, M.C.
Eur. J. Biochem. 94, 375-386, 1979
Bur. J. Biochem. 94, 375-386, 1979
A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat protein of A;Fitle: The primary structure of the coat primary structu
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A;Title: Analysis of the complete nucleotide sequence of the A;Reference number: S01963; MUID:88289362; PMID:3399390
A;Accession: S01964
A;Status: preliminary
                                                                                                                     seeligeriolysin - Listeria seeligeri
C;Species: Listeria seeligeri
C;Date: 22-Nov-1993 #sequence_revision 01-Dec-1995
C;Accession: S22340
Biochim. Biophys. Acta
A;Title: Listeriolysin
A;Reference number: S22
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A; Residues: 1-331
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                                                          R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84,
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C;Superfamily: phage GA coat protein
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1-331 <INO>
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   genes: complete sequence of ilo 2340; MUID:92182018; PMID:1543752
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78.0%;
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Pred. No. 1.7e-44;
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Infect. Immun. 59, 3945-3951, 1991
A;Title: Listeria monocytogenes isolates can be classified A;Reference number: S24230; MUID:92040062; PMID:1937753
A;Accession: S24231
A;Status: nucleic acid sequence not shown; translation not A;Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vysotskaia, V.S.; Yu, G.; Ecker, J.: Theolosubmitted to the EMBL Data Library, February A; Reference number: Z14197
A; Accession: T00646
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C;Accession: T00646
R;Federspiel, N.A.; Palm, C.J.; Conway, A.B.; Kurtz, D.B.; Conway, R;Federspiel, N.A.; Palm, C.J.; Theologis, A.; Davis, R.W.
Vvsotskala, V.S.; Yu, G.; Ecker, J.; Theologis, A.; Davis, R.W.
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A;Molecule type: DNA
A;Residues: 1-530 <ANA>
A;Cross-references: UNIPROT:P31830; EMBL:X60462;
A;Note: the authors translated the codon GCC for C;Superfamily: dipeptide transport protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypothetical protein F316.9 - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 01-Feb-1999 #sequence_revision 01-Feb-1999
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                                                                                                                C;Accession: S24231
R;Rasmussen, O.F.; Beck,
                                                                                                                                                A; Variety: strain 12067
C; Date: 22-Nov-1993 #sequence_revision
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A; Introns: 14/3; 281/3;
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                                                                                                                                                                                                                                             RESULT
                                                                                                                                                                                                        listeriolysin precursor - Listeria monocytogenes
                                                                                                                                                                                         ;Species: Listeria monocytogenes
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33; Conserv
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                                                                                                                T.; Olsen, J.E.;
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24.3%; Pred. No. 1;
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A;Status: preliminary
A;Molecule type: DNA
A;Residues: 413-480 <ME2>
A;Cross-references: GB:M29171
A;Cross-references: GB:M29171
A;Cross-references: GB:M29171
Mol. Microbiol. 4, 2167-2178, 1990
A;Title: Attenuated mutants of the intracellular bacterium Listeria: A;Accession: S12400; MUID:91211627; PMID:1965218
A;Accession: S12400
A;Molecule type: DNA
A;Residues: 483-493 <MIC>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Accession: $05306
A;Molecule type: DNA
A;Residues: 1-529 < COM>
A;Cross-references: EMBL:X15127; NID:g44106; PIDN:CAA33223.1; PID:g44107
A;Cross-references: EMBL:X15127; NID:g44106; PIDN:CAA33223.1; PID:g44107
A;Experimental source: strain EGD
A;Note: this sequence is derived from a weakly hemolytic strain, serotype 1/2a
R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
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R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
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R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cossart, P.
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R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.; Chenevert, P.
R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.; Chenevert, P.; Chenevert, P
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C;Date: 21-Oct-1992 #sequence revision 21-Oct-1992 #text_change 09-Jul-2004
C;Accession: A43505; S05306; A47606; S12400; A61079
R;Mengaud, J.; Vicente, M.F.; Chenevert, J.; Pereira, J.M.; Geoffroy, C.; G
Infect. Immun. 56, 766-772, 1988
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A;Cross-references: UNIDROT:P13128; EMBL:X60035; NID:g44110; PIDN:CAA42639.1; PID:g44117
A;Experimental source: strain 12067, serotype 4b
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1991
A;Experimental source:
C;Genetics:
A;Gene: hlyA; lisA
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Nucleic Acids Res. 17, 6406, 1989
A;Title: Nucleotide sequence of the listeriolysin gene from a Listeria monocytogenes ser A;Reference number: S05306; MUID:89366884; PMID:2505236
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A; Accession: A43505
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A;Reference number: A43505; MUID:88153053; PMID:3126142
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;Keywords: virulence factor
;Keywords: virulence factor
;1-25/Domain: signal sequence #status predicted <NAT>
;26-529/Product: listeriolysin #status predicted <MAT>
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Best Local Similarity
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22.0%; Pred. No. 4
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C;Superfamily: upreprint C;Superfamily: upreprint C;Keywords: virulence factor
F;1-25/Denain: signal sequence #status predicted <SIG>F;1-25/Denain: signal sequence #status predicted <MF;26-529/Product: listeriolysin O #status predicted <MF;26-529/Product: listeriolysin O ---- or F; DB 2;
         RESULT 9
S57537
MXTI protein - yeast (Saccharomyces cerevisiae)
MXTI protein - yeast (Saccharomyces cerevisiae)
N,Alternate names: protein N2302; protein YML085w
C;Species: Saccharomyces cerevisiae
C;Species: Saccharomyces cerevisiae
C;Adte: 10-Oct-1995 #sequence revision 10-Nov-1995 #text_change 09-Jul-2004
C;Accession: S57537; S50279; S63024; S63017; S65096
R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M.
submitted to the EMBL Data Library, June 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Science 294, 849-852, 2001
A; Aunst, F.; Kurapkat, G.; Madueno, E.;
A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.;
ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.;
A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
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R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloecker, p.; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.; Fsihi, H.;
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C;Superfami
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A; Residues: 1-529 <GLA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species:
A;Reference number: S57533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          listeriolysin O precursor [imported] - Listeria monocytogenes (strain EGD-e)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Superfamily: dipeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cross-references: UNIPROT:P13128; GB:NC_003210; PIDN:CAD00729.1; PID:g16409567; GSPDB:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Date: 27-Nov-2001
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                                                                                                                                                                                                                                                                                                                                                                                                         49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13 RDGKQTLVL------VPRGVNPTNGVASLSQAGA------VPALEKRV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Listeria monocytogenes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 RDGKQTLVL-----NPRGVNPTNGVASLSQAGA----
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                                                                                                                                                                                                                                                                                                                                                                                                       TVSVSQPSRNRKWYKVQVKIQNPTACTANGSCDPSVTR-----QKYADVTFSFTQYSTD 102
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22.0%; Pred. No. 4
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Pred. No. 4
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Wehland,
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Overall

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A;Molecule type: DNA
A;Residues: 1-29,'G',31-808,'TMKTCLNYH'.<VER>
A;Residues: 1-29,'G',31-808,'TMKTCLNYH'.<VER>
A;Cross-references: EMBL:U09129; NID:gS20475; PIDN:AAC49470.1;
A;Coler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M.
submitted to the Protein Sequence Database, April 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 569-830 <POE>
A;Cross-references: EMBL:Z71361; MIPS:YNL085w
A;Experimental source: strain S288C
R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha,
Yeast 12, 485-491, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R; Poehlmann, R.; Philippsen, submitted to the Protein Seq
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A;Title: Sequence of MKY1, needed for propagation of M(2)
A;Reference number: S50279; MUID:95176705; PMID:7532890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Accession: S57537
A;Molecule type: DNA
A;Residues: 1-830 <SOL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yeast 12, 485-491, 1975
A;Title: The sequence of a 17 933 bp segment of Saccharomyces
A;Reference number: S65092; MUID:96310628; PMID:8740422
A;Molecule type: DNA
A;Residues: 1-1502 <JOH>
A;Cross-references: EMBL:U20865; NID:g662330; PIDN:AAB67387.1; PID:g662331; GSPDB:GN0001
                                                                                A;Description: The sequence of A;Reference number: S59386 A;Accession: S59400
                                                                                                                                                                                                      N;Alternate names: protein L9672.1; protein Ilakzow; reynamow; protein C;Species: Saccharomyces cerevisiae
C;Date: 30-Sep-1991 #sequence revision 23-Feb-1996 #text_change 12-Nov-1999
C;Accession: S59400; A31312; S15447; S05804; S15446
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C;Superfamily: Saccharomyces cerevisiae MKT1 protein
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                                                                                                                                                                R; Johnson, D. submitted to the EMBL Data Library,
                                                                                                                                                                                                                                                                                                  CYC1/CYP3 transcription activator - yeast (Saccharomyces cerevisiae) N;Alternate names: protein L9672.1; protein YLR256w; regulatory prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: DNA
A;Residues: 1-830 <SOF>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;Cross-references: EMBL:X89016; NID:g887621; PIDN:CAA61425.1;
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;Residues: 1-830 <S
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Vermut, M., Widner, W.R.; Dinman, J.D.; Wickner, R.B.
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Best Local
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                                                                                                                                         <u>د</u>
                                                                                                                                            cerevisiae
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R;Creusot, F.; Verdiere, J.; Gaisne, M.; Slonimski, P.P. J. Mol. Biol. 204, 263-276, 1988
A;Title: CYP1 (HAP1) regulator of oxygen-dependent gene A;Reference number: S15447; MUID:89125585; PMID:2851658
A;Accession: S15447
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R;Pfeifer, K.; Kim, K.S.; Kogan, S.; Guarente, L.
Cell 56, 291-301, 1989
A;Title: Functional dissection and sequence of yeast HAP1 activator
A;Reference number: A31312; MUID:89106221; PMID:2643482
A;Accession: A31312
A;Molecule type: DNA
A;Residues: 1-144,'I',146-322,'R',324-454,'N',456-507,'M',509-586,'I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A,Map position: 12R
C,Superfamily: regulatory protein HAP1; GAL4 zinc binuclear cluster homology
C,Superfamily: heme binding; transcription regulation; zinc finger
C,Keywords: DNA binding #status predicted <DNA>
F,1-149/Domain: DNA binding #status predicted <DNA>
F,59-98/Domain: GAL4 zinc binuclear cluster homology <GAL4>
F,64-84/Region: zinc finger CCC motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   F;177-189/Region: glutamine-rich
F;245-445/Domain: heme binding #status predicted <HEM>
F;245-445/Domain: heme binding #status predicted <F;239-304,323-328,347-352,373-378,389-394,415-420/Region:
F;1308-1481/Domain: activation element #status predicted <F;1388-1481/Region: acidic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Note: the sequence is from mutant CYP1-18 C;Genetics:
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A;Residues: 1-62,'R', 64-1305,'Y',1306-1470,'LVDFYRADFPIWE'
A;Cross-references: EMBL:X13793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 1-144,'I',146-322,'R',324-454,'N',456-507,'M',509-586,'K',588-882,'N',884-9594. A;Cross-references: EMBL:J03152; NID:g171645; PIDN:AAA34662.1; PID:g171646 R;Creusot, F.; Verdiere, J.; Gaisne, M.; Slonimski, P.P.
                                                                                                                                                                                                                                                                                                                                 Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete sequence of ilo
A;Reference number: S22340; MUID:92182018; PMID:1543752
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Gene: SGD:HAP1; CYP1; MIPS:YLR256w
A;Molecule type: DNA
A;Residues: 1-319,'T',321-528 <KRE>
A;Crose-references: EMBL:X60461
C;Genetics:
                                                                                                                         submitted to the EMBL Data A; Reference number: $36683
                                                                                                                                                                                                                                                                                                                                                                                                                 R; Haas, A.; Dumbsky, M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ivanolysin precursor - Listeria ivanovii
                                                                                                         A;Accession: S36683
                                                                                                                                                                                    R;Kreft,
                                                                                                                                                                                                          A; Note: the authors translated the
                                                                                                                                                                                                                                     A; Cross-references: UNIPROT: P31831; EMBL: X60461
                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-528 < HAS>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Species: Listeria ivanovii
;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              54 QPSRNR-KNYKVQVKIQNP----
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35; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            S22341; S36683
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SGD:S0004246; MIPS:YLR256w
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                                                                                                                                                         Library,
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Pred. No. 17
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RESULT 12
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C;Superfamily: dipeptide transport protein
F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-528/Product: ivanolysin #status predicted <MAT>
                                                                                                                                                                            A;Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrobacterium tun
A;Reference number: A97359; MUID:21608551; PMID:11743194
A;Accession: C98221
                                                                                                                                                                                                                                       R;Goodner, B.; Hinkle, G.; Gattung, S.; Miller, N.; Blanchard, M.; Qurollo, A.; Liu, F.; Wollam, C.; Allinger, M.; Doughty, D.; Scott, C.; Lappas, C.; Science 294, 2323-2328, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:P91584; C;Genetics:
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                                                                                                                                                                                                                                                                                                                  hypothetical protein AGR_L_1428 [imported] - Agrobacterium tumefaciens (strain C;Species: Agrobacterium tumefaciens C;Species: Agrobacterium tumefaciens C;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Bird, A.P.; Clark, V.; Jones, S.J.; Leitgeb, S.; Dobson, submitted to the EMBL Data Library, December 1996 A;Reference number: Z21049
A;Accession: T31660
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                                                                                                    A;Cross-references:
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                                                                                                                       ;Molecule type: DNA
;Residues: 1-136 <KUR:
                                                                                                                                                           Status: preliminary
                                                                                                                                                                                                                                                                                                        Accession: C98221;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ETVTLGNIGRDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYK
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                                                                                                  UNIPROT: Q8U8F3; GB: AE007870;
                                          chromosome
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Pred. No. 8.7;
17; Mismatches
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Pred. No. 9
  Score
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  76.5;
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  Length 136;
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                                                                                                PID:g15159127;
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                                                                                                                                                                                                                                                          B.; Goldm
Markelz,
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                                                                                                                                                                                                                                                            Goldman,
kelz, B.
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A;Experimental source: strain C58 (Dupont)
C;Genetics:
A;Gene: Atu4139
                                                                                                                                                                                                                                                                                                                                                                                                                  A;Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58 A;Reference number: AB2577; MUID:21608550; PMID:11743193 A;Accession: AE3065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Authors: Yoo, H.; Tao, ster, E.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; F.; Karp, P.; Romero, P.; Zhang, S. Science 294, 2317-2323, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 C;Accession: AE3065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conserved hypothetical protein Atu4139 [imported] - C;Species: Agrobacterium tumefaciens
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                                                                                                                                                                                                                                                                                                                                                                                                   A;Status:
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                                60 KNYKVQVKIQNPTACTA-----NGSCDPSVTRQKYADVTFSF-TQYSTDEERAFVRTE 111
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                                                                                                                                                                Similarity 29; Conserv
GOYRVKFAIHNLSMTTAAQVNVRGDLEQNGASPENADVTFDYVASESKDNGTLFFRSD 120
                                                                                  ISKDGKQTESADPHWIEWVTGTISTLLVAAMFGWIAYDIYRYSPEEARFEIAVTGVEGQT 62
                                                                                                                       IGRDGKQTLVLNPRGVNPTNGVASLSQAGAV-----PALEKRVTVSVSQPSRNR
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Mismatches
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C;Species: Mycoplasma genitalium
C;Date: 17-Nov-1995 #sequence revision 17-Nov-1995 #text_change 09-Jul-2004
C;Accession: C64221; JQ0092; S18702; S18703
R;Fraser, C.M.; Gocayne, J.D.; White, O.; Adams, M.D.; Clayton, R.A.; Fleischmann, M.; Fraser, C.M.; Mguyen, D.; Utterback, T.R.; Saudek, D.M.; Phillips, C.A.; Merri hypothetical 114K protein (MgPa 3' region) - Mycoplasma genitalium

C.A.; Venter, J.C. Science 270, 397-403, 1995
Science 270, 397-403, 1995
A;Title: The minimal gene complement of Mycoplasma genitalium A;Reference number: A64200; MUID:96026346; PMID:7569993
A;Accession: C64221

A;Status: nucleic acid sequence DNA not shown; translation not shown

A;Residues: 1-1052 <TIGR>
A;Cross_references: UNIPROT:P22747; GB:U39696; GB:L43967; 3. T. NID:g1045869; Hu, P.C PID:g1045877;

A;Experimental source: strain G-37 R;Inamine, J.M.; Loechel, S.; Collier, Gene 82, 259-267, 1989

Mycoplasma

genitalium

ZI.

A;Title: Nucleotide sequence of the MgPa (mgp) operon of A;Reference number: JQ0090; MUID:90060815; PMID:2583522 A;Accession: JQ0092

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A;Residues: 1-1052 <INA>
A;Cross-references: GB:M31431; NID:g150157; PIDN:AAA25421.1; PID:g150160
A;Experimental source: strain G-37 (ATCC 33530)
A;Experimental source: strain G-37 (ATCC 35530)
A;Peterson, S.N.; Schramm, N.; Hu, P.; Bott, K.F.; Hutchison III, C.A.
Nucleic Acids Res. 19, 6027-6031, 1991
A;Title: A random sequencing approach for placing markers on the physical map of Mycopla A;Reference number: $18693; MUID:92051396; PMID:1945886
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A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 902-964 <PET>
A;Cross-references: EMBL:X61525
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, September 1991
A;Note: the nucleotide sequence was submitted to the EMBL Data Library,
Search completed: January 4, 2005, 09:17:29
Job time : 9.03224 secs
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: January 4, 2005, 09:02:16; Search time 36.2676 Seconds (without alignments)
2094.140 Million cell updates/sec

Title: US-10-622-064-6
Perfect score: 664
Sequence: 18LOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1825181 seqs, 575374646 residues
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 Minimum DB seq length: 0
Maximum DB seq length: 200000000

Total number of hits satisfying chosen parameters:

1825181

se : UniProt_02:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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P12351 saccharomyc	Q95ru8 drosophil	Q9v3k8 drosophila	Aam00519 saccharon	Q8tg55 saccharomy	Q8tg52 saccharomy	Q8tg45 saccharomy	Q8tf89 saccharomyc	Q8tf87 saccharomyc	P40850 saccharomyc	Aat03000 listeria	Q6eaj4 listeria	Q6eae0 listeria	Q6eab7 listeria

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Golmohammadi R., "The crystal stru- Structure 4:543-5 -I- FUNCTION: For- This SWISS-PROT e- between the Swiss the European Bioi- use by non-prof modified and this	SEQUENCE. MEDLINE=71288580; Pubmed Maita T., Konigsberg W.; "The amino acid sequence J. Biol. Chem. 246:5003- [5] X-RAY CRYSTALLOGRAPHY (3	REVISIONS TO 1-60 REVISIONS TO 1-60 MEDLINE=77118576; Stoll E., Wilson "Revised amino ac and 60."; J. Biol. Chem. 25	(2) (2) (2) (2) (3) (4) (5) (5) (6) (7) (7) (8) (8) (8) (9) (8) (9) (9) (9) (9) (9) (9) (9) (9) (9) (9	TaxID=1: _TaxID=1: ENCE FROI INE=9410: Ovska T.I V., Pumpo ombinant nbled in	LT 1 BPOBE COAT BPOBE P03615 21-JUL-1986 21-JUL-1986 01-OCT-2004 Coat protein Bacteriophag
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PDB; 1QBE; X-ray; A
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EMBL; V00643; CAA23992.1;
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AAM33126;
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                                              Bacher J.M., Bull J.J., Ellington A.D.;
"Evolution of phage with chemically ambiguous
BMC Evol. Biol. 3:24-24(2003).
EMBL; AY099114; AAM33126.1; -.
                                                                                                      Bacteriophage Q-beta.
Viruses; ssRNA positive-strand viruses, no
Allolevivirus; Allolevivirus subgroup III.
NCBI_TaxID=12009;
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                                Coat protein
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PubMed=14667253;
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HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule ac
InterPro; IPR002703; Levi_coat.
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
PROSITE; PS00030; RRM RNP 1; UNKNOWN 1.
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Kozlovska T.M., Cielens I., Dreilinna D.,
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EMBL; M99039; AAA16663.1; -.
SEQUENCE 329 AA; 36134 MW; E0F30215A8C197CB CRC64;
  Pfam; PF01819; Levi
Coat protein.
SEQUENCE 133 AA;
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MEDLINE=96190948; PubMed=8609616;

MEDLINE=76.190948; PubMed=8609616;

Beekwilder J., Nieuwenhuizen R., Poot R., van Duin J.;

Beekwilder J., Nieuwenhuizen R., Poot R., van Duin J.;

"Secondary structure model for the first three domains

"Secondary structure model for the first three domains

Control of A-protein synthesis.";

J. Mol. Biol. 256:8-19(1996).
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MEDLINE=94109687; PubMed
Kozlovska T.M., Cielens
Ose V., Pumpens P.;
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998) to the EMBL/GenBank/DDBJ d
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GO; GO:005198; F:structural molecule activity;
InterPro; IPR002703; Levi coat.
InterPro; IPR000504; RNA_rec_mot.
Pfam; PF01819; Levi_coat; 1.
PROSITE; PS00030; RRM_RNP 1; UNKNOWN 1.
SEQUENCE 329 AA; 35880 MW; 372B1DD5DDB52F15
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J. Mol. Biol. 247:903-917(1995)
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ
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"Secondary structure model for the first three
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GO; GO:005198; F:structural molecule activity;
InterPro; IPR002703; Levi coat.
InterPro; IPR000504; RNA_rec_mot.
Pfam; PF01819; Levi coat; 1.
PROSITE; BS00030; RRM RNP 1; UNKNOWN 1.
SEQUENCE 329 AA; 35893 MW; 3E33CD821EE625F4
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MEDLING-88289362; PubMed=3399390;
MITTALE PROPERTY OF THE PROPERTY O
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Submitted (MAR-1998) to the E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Beekwilder J., Nieuwenhuizen R., Poot R., van Duin J., "Secondary structure model for the first three domains Control of A-protein synthesis."; J. Mol. Biol. 256:8-19(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=96190948; PubMed=8609616;
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                                                      entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                 This SWISS-PROT entry is copyright. It is produced th between the Swiss Institute of Bioinformatics and the European Bioinformatics Institute. There are no use by non-profit institutions as long as its con
                                                                                                                                                                                                                                                         Hirashima A., Hirose T., 1
"Analysis of the complete coliphage SP.";
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Mol. Biol.
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ative 13;
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to the EMBL/GenBank/DDBJ
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Pred. No. 1.6e-44;
3; Mismatches 14
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Coat protein; RNA-binding.
132 AA; 14129 MW;
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PIR; S01964; S01964.
HSSP; P03615; 1QBE.
InterPro; IPR002703; Levi_coat.
Pfofm; PF01819; Levi_coat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboratio between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are restrictions on it use by non-profit institutions as long as its content is in no was modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic Acids Res. 16:6205-6221(1988).
-I- FUNCTION: Not yet known.
-I- MISCELLANEOUS: The readthrough protein A1 includes the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE-88289362; PubMed=3399390;
MITTOSE T., Inayama S.,
Hirashima A., Hirose T., Inayama S.,
"Analysis of the complete nucleotide
coliphage Sp.";
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01-MAR-1989 (Rel. 10, Last sequence update)
01-MAR-2094 (Rel. 43, Last annotation update)
Readthrough protein Al [Contains: Coat protein].
Bacteriophage SP.
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                                                                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                                                                                                                                                                          331 AA;
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Pred. No. 4.2e-43;
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4A6642E4B52C6582 CRC64;
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Al protein.
Enterobacteria phage NL95.
Viruses; ssRNA positive-strand viruses,
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064310;
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01-AUG-1998
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SEQUENCE 13
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GO; GO:0005198; F:structural molecule activity; IEA.
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ
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Beekwilder J., Nieuwenhuizen R., Poot R.,
"Secondary structure model for the first
Control of A-protein synthesis.";
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01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2003 (TrEMBLrel. 24,
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RESULT 13
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Best Local S
Matches 99
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EMBL; APG59243; AAC14704.1; -
HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule ac:
InterPro; IPR002703; Levi_coat.
InterPro; IPR002703; Levi_coat.
InterPro; IPR00504; RNA rec_mot.
Pfam; P901819; Levi_coat; 1.
Pfam; P901819; Levi_coat; 1.
PSCOUENCE 330 AA; 36175 MW; 961855F40
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P03616;

21-JUL-1986 (Rel. 01, C

21-JUL-1986 (Rel. 01, I

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Submitted (APR-1998) to the
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NCBI_TaxID=75725;
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
Coat protein; Direct protein se
SEQUENCE 131 AA; 14535 MW;
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                                                                                         Eur. J. Biochem. 94:375-386(1979).
-!- FUNCTION: Forms the phage shell;
PIR; A04225; VCBPP1.
                                                                                                                                                                               MEDLINE=79148387; PubMed=107028; Dhaese P., Vandekerckhove J., van I "The primary structure of the coat
                                                                                                                                                                                                                                                                                                                           Viruses; ssRNA positive-strand unclassified Leviviridae.
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  protein sequencing; RNA-binding.
14535 MW; E7E639E1E50FC612 CRC64;
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75.0%; Pred. No. 6.8¢
tive 11; Mismatches
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RESULT 15
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                                           P31830;
01-JUL-1993
01-JUL-1993
05-JUL-2004
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MEDLINE=22350095; PubMed=12461651;
Kiss H., Darai E., Kiss C., Kost-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                GO; GO:0016021; C:integral to membrane;
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Mamm. Genome 13:646-655(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=Tmem7
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Listeria seeligeri.
Bacteria; Firmicutes;
                      Name=1so;
                                Seeligeriolysin precursor
                                                                                                                                                                                                                                                                                                                            Transmembrane.
SEQUENCE '473 AA;
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                   26, Created)
26, Last sequence update)
44, Last annotation update)
recursor (Thiol-activated cytolysin)
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Rodentia;
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32.3%;
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Last annotation updat
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Pred. No. 0.0025;
4; Mismatches 5
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Pred. No. 0.
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Sciurognathi; Muridae; Murinae; Mus
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Mismatches
 Listeriaceae; Listeria.
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MEDINE-92182018; PubMed=1543752;

Haas A., Dumbsky M., Kreft J.;

"Listeriolysin genes: complete sequence of ilo from Listeria ivanovii and of lso from Listeria seeligeri.";

Biochim. Biophys. Acta 1130:81-84(1:92).

-1- FUNCTION: Sulfhydryl-activated toxin. Is able to lyse cholesterol containing membranes. Can be reversibly inactivated by oxidation. Cholesterol is the receptor for the binding of these toxins to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; X60462; CAA42996.1; -.
prir; S22340; S22340.
HSSP; P19995; 1PFO.
InterPro; IRR001869; Thiol cytolysin.
Pfam; PF01289; Thiol cytolysin; 1.
PRINTS; PR01400; TACTYTOLYSIN.
ProDom; PD007062; Thiol cytolysin; 1.
PROSITE; PS00481; THIOL CYTOLYSINS; 1.
CYtolysis; Hemolysis; Lipid-binding; Signal; Toxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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-I- SIMILARITY: Belongs to the thiol-activated cytolysin family.
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    209
                                                                                                                                                   154 TLSVDLPGMTKKONKIFVK--NPTKSNVNNAVNTLVERWNDKYSKAYPNINAKIDYS--- 208
                                                                         99 YSTDEERAFVRTELAA 114
                                                                                                                                                                                                                                 49 TVSVSQPSRNRKNYKVQVKIQNPTACTANGSCDPSVTR--QKY-----ADVTFSFTQ 98
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26 530 Seeligeriolysin.
485 485 Binding to cholesterol (By similarity).
530 AA; 59181 MW; 416F7A4DD2029866 CRC64;
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Search completed: January 4, 2005, 09:16:15
Job time: 37.2676 secs

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Result
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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ALIGNMENTS

RESULT 1 ABG94278 IID ABG94278 AC ABG9 XX ABG9 XX ABG9 XX ABG9 XX ABG9 XX Huma XX Yacc WPI; 19-JAN-2001; 2001US-0262379P 04-MAY-2001; 2001US-0288549P 05-OCT-2001; 2001US-0326998P 07-NOV-2001; 2001US-0331045P Renner WA, Piossek C; 21-JAN-2002; 2002WO-IB000166 25-JUL-2002 WO200256905-A2 Bacteriophage. cytostatic; antiviral; vaccine; infectious dis (CYTO-) 2002-627351/67. CYTOS BIOTECHNOLOGY AG Bachmann al; antidiabetic; hypoglycaemic; disease. ž Tissot Þ Maurer ۵ Lechner antigen array; Ή, Sebbel

Human; mouse; rat; antimicrobial; antiallergic;

immunomodulatory;

Bacteriophage Q beta coat protein

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06-AUG-2003 10-DEC-2002 ABG94278;

(revised)
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ABG94278 standard;

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Claim diseases. Molecular

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This invention relates to a novel ordered and repetitive antigen array used in the production of vaccines for infectious diseases. The invention also discloses a composition comprising a non-natural molecular scaffold comprising a core particle selected from a core particle of a non-natural origin and a core particle of natural origin and an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond. Also disclosed is an

antigen or antigenic determinant with at least one second

attachment

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RESULT 2
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                                                                                                          Molecular antigen array; vaccine; antigen; antimicrobial; molecular scaffold; amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgB-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus erythematosus; osteoporosis; inflammatory immune disease; myasthenia gravis; multiple sclerosis; immunoproliferative disease; myphadenopathy; Alzheimer's disease; angioimmunoproliferative lymphadenopathy; immunoblastive lymphadenopathy; rheumatoid arthritis; diabetes; infectious disease; factor Xa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antiviral, antidiabetic, or hypoglycaemic activities and may be used in immunisation and as a vaccine. The present sequence represents a protein sequence used to create the compositions of the invention. (Updated on 0-AUG-2003 to correct OS field.)
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MAKLETVILGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR

Query Match Best Local S Matches 329

Similarity

Score 1749; DB 5; Pred. No. 3e-171; ; Mismatches 0;

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CC molecular scaffold comprising: (i) a core particle selected from: (1) a core particle of a non-natural cc core particle of a non-natural origin; and (2) a core particle of natural cc corigin; and (ii) an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least cone second attachment site, where the antigen or antigenic determinant with at least cone second attachment site, where the antigen or antigenic determinant is attachment is selected from: (i) an attachment, and where the second attachment site not naturally occurring with the antigen or antigenic determinant is through at least one non-peptide bond to the first attachment site; and through at least one non-peptide bond to the first attachment site; and condeterminant, where the second attachment site is capable of association conditions included is a process for producing a non-naturally occurring with the antigen or antigenic determinant and the scaffold interact through the association to form an ordered and repetitive antigen array. The composition is used in immunisation and as a vaccine for diseases such as influenza, graft respiratory distress syndrome (ARDS). Crohn's disease, allergic asthma, acut lymphoblastic leukaemia, non-Hodgkin's lymphoma, Grave's disease, gravis, immunoproliferative disease lymphadenopathy, anaphylaxis, adult creaminatoid arthritis, diabetes, multiple sclerosis, Alzheimer's disease, costeoporosis and infectious diseases. The antigens are modified to posses a cleavage site (enterokinase or factor Xa) and a Cysteiner-containing N-creaminal linker peptide which serves as the attachment point to a creaminal linker peptide which serves as the attachment containing N-creaminal linker peptide which serves as the attachment containing N-creaminal linker peptide which serves as the attachment point to the creaminal serves 
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04-MAY-2001;
05-OCT-2001;
07-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 393-394; 418pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Molecular antigen array used in
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Renner WA,
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                                    virus like particle or bacterial protein (the scaffold protein). The present sequence is bacterial protein or peptide which is coupled to the modified antigen to form the molecular antigen array. (Updated on 29-AUG-2003 to standardise OS field)
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LUEOEND R.
STAUFENBIEL
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; 2001US-0326998P.
; 2001US-0331045P.
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                                                                                                                                                                                                                                                                                                                                    New angiotensin peptide moiety carrier conjugate comprising a carrier and an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteriophage Qbeta; coat protein; A1; hypotensive; cerebroprotective; cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction; congestive heart failure; kidney failure; retinal haemorrhage.
                    The invention describes an angiotensin peptide moiety carrier conjugate comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension,
                                                                                                                                                                                                                                                        Claim 16; Page 92-93;
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21-JAN-2002;
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21-JAN-2002; 2002WO-1B000166
08-JUL-2002; 2002US-03937259
18-JUL-2002; 2002US-0396590P
                                                                                                                                                                                                                                                    vaccine composition; virus-like particle; core particle; first attachment site; antigen; antigenic determinant; prion protein; PrP; PrP peptide; vaccine; neuroprotective; antiinflammatory; prion disease; Bovine Spongiform Encephalopathy; BSE; Creutzfeldt-Jakob Disease; coat protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inaemorrhage. The conjugate is also useful for inducing including producing antibodies. This is the amino acid bacteriophage Obeta coat protein Al used in the prepara vaccine conjugates of the invention
                                                                                                                                                                                  24-JUL-2003.
                                                                                                                                                                                                         WO2003059386-A2.
                                                                                                                                                                                                                             Bacteriophage Qbeta.
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Pred. No. 3e-171;
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A vaccine composition for preventing Creutzfeldt-Jakob Disease) comprises

or treating parties a virus-like

prion diseases e particle (e.g.

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Best Local Similarity
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like or a core particle with at least one first attachment site and at
least one antigen or antigenic determinant that is a prion protein (PP)
or its dimer, or a PrP peptide, the antigen or antigenic determinant
being bound to the virus-like or core particle. The vaccine of the
invention may have neuroprotective or antinflammatory activity. The
composition is useful as a medicament or in manufacturing a medicament
for the treatment or prevention of prion diseases. The prion diseases may
include Bovine Spongiform Encephalopathy (BSE) or Creutzfeldt-Jakob
Disease. The present sequence is the amino acid sequence of a coat
protein from a bacteriophage which may be used during the creation of the
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  07-NOV-2001; 2001US-0331045P
18-JAN-2002; 2002US-00050902
21-JAN-2002; 2002WO-IB000166
                                                                                                                                            Unidentified
                                                                                                                                                                      osteopathic;
bone disease;
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                                                         07-NOV-2002; 2002WO-EP012449
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ilarity 100.0%;
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encephalopathy; immune
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Pred. No. 3e-171;
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                                                                                                                                                                      antigenic determinant; RANKL;
system stimulation.
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antiallergic; antiasthmatic; cytostatic; vaccine; virus-like interleukin; IL-5; IL-13; eotaxin; repetitive antigen array; allergic eosinophilic disease; asthma; Hodgkin's lymphoma.

Virus-like particle repetitive antigen array peptide

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                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 329;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 329 AA;
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                                                                                            LATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP
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SSGGAIPFDFTKFDKTKCPIQAVIVVPRA
                                                                                                                                              PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a composition comprising a virus-like particle CC and at least one antigen, which is a protein or peptide of interleukin CC (II)-5, II-13 or eotaxin and is bound to the virus-like particle, or a CC core particle with at least one first attachment site and at least one at CC protein or peptide of II-5, II-13 or eotaxin. The second attachment site one cC protein or peptide of II-5, II-13 or eotaxin. The second attachment site is a nattachment site naturally or not naturally occurring with the CC is an attachment site naturally or not naturally occurring with the CC antigen or antigenic determinant. The second attachment site is capable of association to the first attachment site, and where the antigen or CC antigenic determinant and the core particle interact through the CC association to form an ordered and repetitive antigen array. The CC compositions are useful as medicaments, or for manufacturing a medicament CC or a vaccine for treating allergic eosinophilic diseases, e.g. asthma, or CC Hodgkin's lymphoma and related diseases. This sequence is used to cycles.
                                                                                                                                                                                                                                                                                                                                                                                                       Matches 329;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Composition comprising an ordered and repetitive antigen or antigenic determinant array, useful as a medicament, or for manufacturing a medicament for treating allergic eosinophilic diseases, e.g. asthma, (Hodgkin's lymphoma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 329 AA;
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21-JAN-2002;
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; 2002US-00050902.
; 2002WO-IB000166.
; 2002US-0396636P.
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Pred. No. 3e-171;
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                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                          an animal comprising a virus-like particle, an immunostimulatory substance bound to the virus-like particle, and an antigen mixed with the virus-like particle. The composition or the vaccine is useful in the manufacture of a pharmaceutical for the treatment of a disorder or disease such as allergies, tumours, chronic diseases and chronic viral diseases. The composition is also useful for enhancing an immune response in an animal. This is the amino acid sequence of a bacteriophage Obeta coat protein API protein subunit a virus like particle (VLP) that can be used in the adjuvant of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New compositions comprising a virus-like particle (VLP), an immunostimulatory substance bound to the VLP, and an antigen mixed with the VLP, useful for enhancing immune response or for treating e.g. tumos or chronic viral diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiallergic; cytostatic; virucide; immunostimulant; vaccine;
immune response; virus-like particle; immunostimulatory; allergy; tumour;
chronic disease; chronic viral disease; bacteriophage Qbeta;
                                                                                                                                                                                                                                                                                                                 Sequence 329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention describes a composition for enhancing an immune response in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 2; 252pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bacteriophage Qbeta
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                                                                               LLIDAIDQLNPAYWTLLIAGGGGGSKPDPVIPDPPIDPPPGTGKYTCPFAIWSLEEVYEP 180
                      PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTPRGCRGNGYIDLDATY
                                                                                                                         KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASP
                                                                                                                                           KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDBERAFVRTELAALLASP 120
                                                                                                                                                                                       MAKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR
                                                                                                                                                                                                           MAKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR
                                                                                                                                                                                                                                                  100.0%; Score 1749; DB 8 llarity 100.0%; Pred. No. 3e-171; Conservative 0; Mismatches 0
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LATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP

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RESULT 7
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antigenic array.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADJ67147 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                        or antigenic determinant, where the second attachment site is capable of association to the first attachment site, and where the ghrelin or a ghrelin peptide and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful for treating obesity. The repetitive array may form part of a phage or bacterial display array. This peptide corresponds to a Bacteriophage obeta coat protein which can used as part of the repetitive or antigenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a new composition comprising: (i) a core particle with at least one first attachment site; and (ii) at least one antigen or antigenic determinant with at least one second attachment site, where the antigen or antigenic determinant is ghrelin or a ghrelin peptide, and where the second attachment site being consisting of an attachment site not naturally occurring with the antigen or antigenic determinant and an attachment site naturally occurring with the antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New composition comprising a core particle having a first attachment and an antigen or antigenic determinant which is a ghrelin or ghrelin peptide having a second attachment site, useful for treating obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bachmann MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteriophage Qbeta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bacteriophage Qbeta coat protein for antigen display array.
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                                                                                                                                                                                                                                                                                                                                        Sequence
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hes 329;
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Similarity 100.0%;
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                                KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASP
                                                                                                          KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASP
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                                                                                                                                                                                                                                                        Score 1749; DB 8 Pred. No. 3e-171;
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RESULT 9
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PR 10-J
PR 115-M
XXX (CVT)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neuroprotective; nootropic; vaccine; amyloid beta 1-6 peptide;
core particle; antigen array; Alzheimer's disease; RNA bacteriophage;
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15-MAY-2003;
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Composition comprising a core and an antigenic amyloid beta such as Alzheimer's disease. particle with 1-6 peptide. " useful at least one attachment site, useful for treating diseases

Claim 12; SEQ ID NO 5; 184pp; English.

particle with at least one attachment site, and an antigenic amyloid beta CC one first attachment site; and at least come particle with at least CC one first attachment site; and at least one antigen or antigenic determinant with at least one second attachment site, where the antigen CC or antigenic determinant is a Amyloid beta 1-6 peptide, and where the second attachment site not naturally CC occurring with the antigen or antigenic determinant; or an attachment CC site naturally occurring with the antigen or antigenic determinant; or an attachment CC site and the beta 1-6 peptide and the core particle interact through the cassociation to form an ordered and repetitive antigen array. The CC escendiation is useful for the manufacture of a medicament for treating CC Alzheimer's disease and related diseases. This is the amino acid sequence of an RNA bacteriophage coat protein that can be used in the preparation of the composition and vaccines of the invention.

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ABRSGUAT 10
ABRS640
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antithyroid; antidiabetic; neuroprotective; nootropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis; Alzheimer's disease; osteoporosis; rheumatoid arthritis;
                                               A composition, useful for enhancing an immune response against an antigen or a virus-like particle, enhancing anti-viral protection in an animal, or immunizing or treating tumors or infectious diseases, e.g. viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacteriophage Qbeta.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; virucide; antibacterial; antiparasitic; fungicide;
antiallergic; immunosuppressive; antiaddictive; antiinflammatory;
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28-JUL-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammatory autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antigen presenting cell; APC; immune response; virus like particle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteriophage Q-beta coat protein SEQ ID NO:11.
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                                                                                                                                                                                                                             Storni T,
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                                                                                                                                                             The present invention describes a composition (C) for enhancing an immune cresponse against an antigen or a virus-like particle in an animal. (C) CC comprises a virus-like particle (VLP) bound to at least one antigen, or a CC VLP capable of being recognised by the immune system of the animal. Also described: (1) enhancing an immune response against an antigen or a VLP (C) in an animal comprising introducing (C) into the animal, (2) vaccines (2) comprising (C) together with a pharmaceutical diluent, carrier or (2) cercipient; (3) immunising or treating an animal comprising administering the vaccine; and (4) enhancing anti-viral (C) the animal by administering the vaccine; and (4) enhancing anti-viral (C) protection in an animal comprising introducing (C) into the animal. (C) (C) has cytostatic, virucide, antibacterial, antiparasitic, fungicide, antiblergic, immunosuppressive, antiaddictive, antiinflammatory, (C) antithyroid, antidabetic, neuroprotective, nootropic, osteopathic, etc. (C) (C) antithyroid, antidabetic, neuroprotective, nootropic, osteopathic, etc. (C) (C) antirheumatic and antiarthritic activities. (C) or the vaccines can be (C) used for enhancing anti-viral protection in an animal, or immunising or treating tumours and infections. The vaccine such as viral, bacterial, (C) parasitic or fungal infections. The vaccine compositions are also useful (C) parasitic or fungal infections. The vaccine compositions are also useful (C) autoimmune disease, accesses, disease, diabetes, multiple sclerosis, (C) Alzheimer's disease, accesses, of the massing of infections to disease, diabetes, multiple sclerosis, (C) Alzheimer's disease, accesses, of the present invention. (C) parasition of the present invention. (C) proposent sequences used in the exemplification of the present invention.
Sequence 328
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片 Ś 밁 S 밁 Ś 맑 Ś 멍 Ş Query Match Best Local (Matches 301 240 180 181 121 121 61 52 328; 1 MAKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR Similarity LLIDAIDQLNPAYWTLLIAGGGSGSKPDPVIPDPPDDPPDGTGKYTCPFAIWSLEEVYEP 180 SSGGAIPFDFTKFDKTKCPIQAVIVVPRA .329 KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASP 120 MAKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR SSGGAIPFDFTKFDKTKCPIQAVIVVPRA LATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASP <u>LATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP</u> PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY LLIDAIDQLNPAYW-LLIAGGGSGSKPDPVIPDPPIDPPPGTGKYTCPFAIWSLEEVYEP Conservative 99.1%; 99.7%; <u>,</u> Score 1733.5; Pred. No. 1.2 Mismatches 1.2e-169; B 0, 6 Indels Length 328; 1; 300 240 120 60 239 179 299

ABR44543; ABR44543 standard; protein; 328 ð

Bacteriophage Q-beta coat protein SEQ ID NO:11.

23-OCT-2003 25-JUL-2003

(revised) (first en

entry)

Immunostimulatory; virus-like particle; bacteriophage; HBV; LCMV; hepatitis B virus; lymphocytic choriomeningitis virus; vaccine; immunostimulant; cytostatic; antiallergic; virucide; antibacteria antibacterial;

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the points in all all marks. (c) comprises a compositional actory substance. The immunostimulatory substance is bound to the VLP Also described: (1) enhancing an immune response in an animal by introducing (C) into the animal; (2) producing (C) for enhancing an immune response in an animal by immune response in an animal (2) producing (C) for enhancing an immune response in an animal (3) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; and (4) immunising or treating an animal by: (a) administering the vaccine to the animal; (b) creating a T cell response in the animal by administering the vaccine; or (c) boosting a T cell response in the animal by administering the vaccine; or concine. (C) has immunostimulant, cytostatic, antiallergic, virucide and antibacterial activities. (I) can be used in vaccines for enhancing an immune response in an animal, particularly a mammal or human. CC specifically, (C) is useful for enhancing a mammal or human. The comprising (C) can also be used for immunising or treating an animal, ce.g. humans, sheep, horses, cattle, pigs, dogs, cats, birds, ce.g. humans, sheep, horses, cattle, pigs, dogs, cats, birds, creptiles or fish. (C) is particularly useful in prophylactic or therapeutic vaccines against allergies, tumours (e.g. breast cancers, neuroblastoma, or leukaemia), viral diseases (e.g. influenza, hepatitis, compresent sequences used in the exemplification of the present invention. (Updated on 23-0CT-2003 to standardise OS field)
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New compositions comprising immunostimulatory substances packaged into virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, tumors
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(TISS/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 247-249; 322pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CIEL/)
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                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                     Local Similarity
nes 328; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2003-354564/33
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MEIJERINK E.
LIPOWSKY G.
PUMPENS P.
CIELENS I.
RENHOFA R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CYTOS BIOTECHNOLOGY AG
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                                                                                           MAKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in an animal.
                                                        MAKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR
KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissot A,
Cielens I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention describes a composition (C) for enhancing an im
an animal. (C) comprises a virus-like particle (VLP), and
latory substance. The immunostimulatory substance is bound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001US-0318994P.
2002US-0374145P.
                                                                                                                                                                        Conservative
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                                                                                                                                                                     0;
                                                                                                                                                                                                  Score 1733.5; DB 6; Pred. No. 1.2e-169;
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, Bachmann MF,
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RESULT 12
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response against an antigen or a virus-like particle in an animal. (C) comprises a virus-like particle (VLP) bound to at least one antigen, or a VLP capable of being recognised by the immune system of the animal. Also described: (1) enhancing an immune response against an antigen or a VLP in an animal comprising introducing (C) into the animal; (2) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; (3) immunising or treating an animal comprising administering the vaccine to the animal, or priming or boosting a T cell response in the animal by administering the vaccine; and (4) enhancing anti-viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              graft-versus-host disease; Grave's disease; diabetes; mu
Alzheimer's disease; osteoporosis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cytostatic; virucide; antibacterial; antiparasitic; fungicide; cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory; antithyroid; antidiabetic; neuroprotective; nootropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteriophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteriophage SP coat protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            or a virus-ing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A composition, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bachmann MF,
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                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 173-174;
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                                                                                                                                                                                                                                                                                                 The present invention describes a composition (C) for enhancing an immune
                                                                                                                                                                                                                                                                                                                                                                                                                                     infections
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         presenting cell; APC; immune response; virus like particle;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful for enhancing an immune response against an antigen particle, enhancing anti-viral protection in an animal, r treating tumors or infectious diseases, e.g. viral
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                                                                                                                                                                                                                                                                                                                                                                Immunostimulatory; virus-like particle; bacteriophage; HBV; LCMV; hepatitis B virus; lymphocytic choriomeningitis virus; vaccine; immunostimulant; cytostatic; antiallergic; virucide; antibacterial; immune response; immunisation; allergy; tumour; breast cancer; neuroblastoma; leukaemia; viral disease; influenza; hepatitis; measles;
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   14-SEP-2001;
22-APR-2002;
                                                                                             16-SEP-2002; 2002WO-IB004132
                                                                                                                                                                                                                                                                               Bacteriophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacteriophage SP coat protein SEQ ID NO:16.
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                                                                                                                                                                                                                                                                                                                                             pox; bacterial infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; protein;
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2002US-0374145P.
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Pred. No. 9.9e-92;
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                                                                                                                                                                                                                                                                                                                                             tuberculosis;
                                                                                                                                                                                                                                                                                                                                         pneumonia; syphilis.
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122 LIDAIDQLNPAYW-TLLIAGGGSGSKPD----PVIPDPPIDPPPGTGKYTCPFAIWSLEE

177 176 NFKVQIKLQNPTACTRD-ACDPSVTRSAFADVTLSFTSYSTDEERALIRTELAALLADPL

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61 62 Best Local Sim Matches 196; Query Match

Similarity

58.3%;

Conservative

45;

Score 978; DB 6; Length Pred. No. 9.9e-92; 5; Mismatches 77; Indel

Indels

18;

Gaps

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AKLETYTIGNIGKDGKQTIVI.NPRGVNPTNGVASISQAGAVPALEKRVTVSVSQPSRNRK 61

AKLNQVTLSKIGKNGDQTLTLTPRGVNPTNGVASLSEAGAVPALEKRVTVSVAQPSRNRK

121

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δ 밁 S 밁

234

177 120

VYEPPTKNRPWPIYNAVBLQPREFDVALKDLLGNTKWRDWDSRLS---YTTFRGCRGNGY IVDAIDNLNPAYWAALLVASSGGGDNPSDPDVPVVPD--VKPPDGTGRYKCPFACYRLGS

VALEDFLGNTNWRNWDQRLSDYDIANRRCRGNGY

IDLDATYLATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSI--NAYCSLSDIAAYHADG

291 236 233

IDLDATAMOSDDFVLSGRYGVRKVKFPGAFGSIK---YLLNIOGDAWLDLSEVTAYRSYG

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The present invention describes a composition (C) for enhancing an immune response in an animal. (C) comprises a virus-like particle (VLP), and an C immunostimulatory substance is bound to the VLP. Also described: (1) enhancing an immune response in an animal by CC introducing (C) into the animal; (2) producing (C) for enhancing an CC pharmaceutical diluent, carrier or excipient; and (4) immunising or creating an animal by: (a) administering the vaccine to the animal; (b) priming a T cell response in the animal by administering the vaccine; or CC (c) boosting a T cell response in the animal by administering the vaccine; or CC (c) boosting a T cell response in the animal by administering the vaccine; or CC (c) boosting a T cell response in the animal by administering the vaccine; or CC (c) bas immunostimulant, cytostatic, antiallergic, virucide and immune response in an animal, particularly a mammal or human. CC specifically, (C) is useful for enhancing a B cell response, a T cell response, or a cytotoxic T-lymphocyte (CTL) response. (C) or a vaccine CC comprising (C) can also be used for immunising or treating an animal, CC e.g., humans, sheep, horses, cattle, pigs, dogs, cats, rats, birds, cc therapeutic vaccines against allergies, tumours (e.g. breast cancers, CC neuroblastoma, or leukaemia), viral diseases (e.g. influenza, hepatitis, comeasles or chicken pox), or bacterial infections (e.g. tuberculosis, particularly and animal and particularis (e.g. tuberculosis, compresent sequences used in the exemplification of the present invention of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Maurer P, L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New compositions comprising immunostimulatory substances packaged into virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, tumors
Sequence 329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 252-253; 322pp;
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(MAUR/)
(TISS/)
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LIPOWSKY G.
PUMPENS P.
CIELENS I.
RENHOFA R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SCHWARZ K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Renhofa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ĀĢ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
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VIVGFWRDPSSGGAIPFDFTKFDKTKCPIQAVIVVP

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ADD24123
ID ADD24
XX ADD244
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KW First
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KW PrP;
KW PrP;
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KW PrD
N W0200
XX Creut
XX Bacte
PN W0200
XX Hack
PN W17-JF
PR 18-JF
PR 08-JF
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                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to a novel vaccine composition comprising a virus-
like or a core particle with at least one first attachment site and at
least one antigen or antigenic determinant that is a prion protein (PrP)
or its dimer, or a PrP peptide, the antigen or antigenic determinant
being bound to the virus-like or core particle. The vaccine of the
invention may have neuroprotective or antificaturing a medicament
for the treatment or prevention of prion diseases. The prion diseases may
include Bovine Spongiform Encephalopathy (BSE) or Creutzfeldt-Jakob
Disease. The present sequence is the amino acid sequence of a coat
protein from a bacteriophage which may be used during the creation of the
vaccine composition of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A vaccine composition for preventing or treating prion diseases Creutzfeldt-Jakob Disease) comprises a virus-like particle (e.g. phage) and at least one prion protein or peptide bound to the vi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JAN-2002; 2002WO-IB000166.
08-JUL-2002; 2002US-0393725P.
18-JUL-2002; 2002US-0396590P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JAN-2003; 2003WO-EP000460
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                                                                                                                                                                                                                                                                                                                                                 Sequence 329
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                                 NFKVQIKLQNPTACTRD-ACDPSVTRSAFADVTLSFTSYSTDEERALIRTELAALLADPL
                                                                                                                     AKLNQVTLSKIGKNGDQTLTLTPRGVNPTNGVASLSEAGAVPALEKRVTVSVAQPSRNRK
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                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                Score 978;
Pred. No. 9.
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e.g. RNA-
e virus-like
                                                                                                                                                                                                                                     Gaps
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                                                    The invention relates to a composition comprising a core particle having at least one first attachment site, and at least one antigen or antigenic determinant having at least one second attachment site. The antigen or antigenic determinant is a RANKL protein, RANKL fragment or RANKL peptide. The second attachment site is (non-) naturally occurring with the antigen or antigenic determinant, and is capable of association to the first attachment site. The antigen or antigenic determinant and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful as a medicament, or composition is especially useful for as a vaccine for therapy or prophylaxis of bone diseases, particularly mammalian encephalopathies, and for stimulating mammalian immune system. This sequence represents a protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compositions comprising a core particle and at least one antigen antigenic determinant, useful for as a vaccine for therapy or prophyl of bone diseases, particularly mammalian encephalopathies.
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18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002UO-1B000166.
19-JUL-2002; 2002US-0396635P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   osteopathic; vaccine; core particle;
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Search completed: January 4, 2005, 09:10:59 Job time : 99.9054 secs	62 NYKVQVKIQNPTACTANGSCDPSVTROAYADVTFSFTQYSTDEERAFVRTELAALLASPL 121	55.9%; Score 978; DB 7; Length 329; Similarity 58.3%; Pred. No. 9.9e-92; 6; Conservative 45; Mismatches 77; Indels 18; Gaps AKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK AKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK AKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK

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US-09-136-574A-43
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Patent No. 6294366
GENERAL INFORMATION:
APPLICANT: Farrington, Graham 1
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INFORMATION FOR SEQ ID NO: 43:
                                                                                                                                                                                             COUNTRY: FA
COUNTRY: USA
ZIP: 19477
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/136,574A
FILING DATE: 19-Aug-1998
CIASSIFICATION CURROWN>
PRIOR APPLICATION NUMBER: US 08/932,571
FILING DATE: September 19, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
REFERENCE/DOCKET NUMBER: 1997US001/CI
TELECOMMUNICATION INFORMATION:
 TOPOLOGY: linear MOLECULE TYPE: No. 6294366e SEQUENCE DESCRIPTION: SEQ ID NO: 43:
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CORRESPONDENCE ADDRESS:
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                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 1426 amino a
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STATE: PA
                                                      STRANDEDNESS: single
                                                                           TYPE: amino acid
                                                                                                                                                                 TELEPHONE: 215-540-9200
TELEFAX: 215-540-5818
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Bergquist, Peter
Daniels, Roy
Morgan, Hugh W.
Williams, Diane P.
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Treating Cellulose Containing Fa
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 17932
LENGTH: 1326
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION: APPLICANT: Marc J.
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                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/074,788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                      LENGTH: 13
TYPE: PRT
                                                                                                                                                                                                                                                                                                                  ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    517 VRFNKNDWSNYNQADDWSWLQSMTDYGENAKVTLY-VDGVLV--WGQEP--GGATPAPTA 571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            457 VDGDKPQSAVCDWAQIGASNVTFNFVKLSSGVSGADYYLEVGFSSGAGQLQPGKDAGDIQ 516
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 453
                                  123 --IDAIDQLNPA-----YWTLL--IAGGGSGSKPDPV---IPDPPIDPPPG------
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     80;
                                                                                                                                                                                17 KOTLVLNPRG----VNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQ-VKIQN 71
                                                                                                                                                                                                                         92;
                                                                       VSQASANQRKGRCGRVEPGICVRLYSEEDFNARPAFTDPE--ILRTNLAAVILQMLHLRL
                                                                                                          PTACTAN----GSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPLL---
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GDIEAFPFIEPPDGKAIKDGFTLLQELSAVNREGQLTPLGRQLARLPIDPRLGRMLLEAA 512
                                                                                                                                               EQQKIFQPRPGRKIVLSTN-VAETSL--TVPGIRYVIDSGTARISRYSYRAKVQRLPIEA
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VENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
VENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
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                                                                                                                                                                                                                         48; Mismatches
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US-08-289-222E-3
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                                                                                                                TELEFAX: 202/638-4810
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20005-5701
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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FILING DATE: 12-AUG-
PRIOR APPLICATION DATA:
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   TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                 SEQUENCE CHARACTERISTICS
                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: KITTS, MONICA CHIN
                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
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                                      STRANDEDNESS:
                                                          TYPE: amino acid
                                                                                                                                                                                          NAME: KITTS, MONICA CHIN REGISTRATION NUMBER: 36,105 REFERENCE/DOCKET NUMBER: P5
                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: DE P
FILING DATE: 07-JUL-1994
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                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
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655 FIFTEENTH STREET, N. W., G STREET LOBBY,
SUITE 330
                                                                           401 amino acids
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Query Match
Best Local Similarity
Tatches 77; Conserv
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                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 20005-5701
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/054,526B
FILING DATE: 03-APR-1998
                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 44 23 190.3
FILING DATE: 01-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EPO 92102324.8
APPLICATION NUMBER: PCT/
FILING DATE: 12-FEB-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 12-AUG-1994
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                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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BECHTOLD, ROLF
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21.7%;
                                              PCT/EP93/00350
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Pred. No. 0.
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W., G STREET LOBBY,
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SEQUENCE CHARACTERISTICS:
                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Neidhardt, Helge
APPLICANT: Paulista, Michael
TITLE OF INVENTION: NEW GROWTH/DIFFERENTIATING
TITLE OF INVENTION: THE TGF- FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: P5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/638-5000
               PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P
FILING DATE: 10-AUG-1993
                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
PRIOR APPLICATION DATA:
                                                                     APPLICATION NUMBER: US/0 FILING DATE: 10-AUG-1994 CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                            STATE: D.C.
                                                                                                                                                                                                                                                                                           CITY: Washington
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21.7%; Pred. No. 0.18;
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th Street N.W. Suite
                                                                                                     US/08/288,508C
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                                                                                                                                             Version
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; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-288-508C-2
                                                                                                                                                                                                                                                                                                                                                                                                    US-08-981-490B-1
               PRIOR FILING DATE: 1995-07-12
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 501
                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/08981490B Patent No. 6531450
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                    APPLICANT: Bechtold, Rolf
APPLICANT: Bechtold, Rolf
APPLICANT: Paulista, Michael
APPLICANT: Unsicker, Klaus
TITLE OF INVENTION: USE OF MP52 OR MP121 FOR TREATING AND PREVENTING DISEASES OF TITLE OF INVENTION: MERVOUS SYSTEM
FILE REFERENCE: 100564-07032
CURRENT APPLICATION NUMBER: US/08/981,490B
CURRENT FILING DATE: 1998-05-18
PRIOR APPLICATION NUMBER: PCT/EP96/03065
PRIOR APPLICATION NUMBER: DS/195 25 416.3
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LENGTH: 501 amino acid
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REGISTRATION NUMBER: P-41,0
REFERENCE/DOCKET NUMBER: P5.
TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                           Pohl, Jens
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; ORGANISM: Homo sapiens US-08-981-490B-1
                                                                                                                                                                                                                                                        US-09-386-450D-2
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                                                                                                                                                    Query Match
Best Local Similarity
Matches 77; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: DE P 44 20 157.5
PRIOR FILING DATE: 1994-06-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 08/288,508
PRIOR FILING DATE: 1994-08-10
PRIOR APPLICATION NUMBER: DE P 43 26 829.3
PRIOR FILING DATE: 1993-08-10
PRIOR APPLICATION NUMBER: DE P 44 18 222.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 100564-09022
CURRENT APPLICATION NUMBER: US/09/386,450D
CURRENT FILING DATE: 1999-08-31
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APPLICANT: Neidhardt, Helge
APPLICANT: Paulista, Michael
TITLE OF INVENTION: NEW GROWTH/DIFFERENTIATING FACTOR OF TGF-? Family
FILE REFERENCE: 100564-09022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 41
SOFTWARE: DALLEY OF SEQ ID NOS: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1994-05-25
                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: DOMAIN
LOCATION: (1)...(501)
OTHER INFORMATION: TGF-beta protein MP-52 precursor
                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 501
TYPE: PRT
ORGANISM: Homo sapiens
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nes 77; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               122 VTPKGQLP--GGKAPPKAGSVPSSFLLKKAREPGPPREPKEPFRPPPITPHEYMLSLYRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2, Application US/09386450D
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                                                    VTPKGQLP--GGKAPPKAGSVPSSFLLKKAREPGPPREPKEPFRPPPITPHEYMLSLYRT
                                                                                                   LNPRGVNPTNGVASLSQAGAVP-----ALEKRVTVSVSQPSR-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GKKPGAFGNIERFIYLKSINAYCSLS------DIAAYHADGV
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                                                                                                                                                       Conservative
-NRKNYKVQVKIQNPTACTANGSCD------PSVTRQAYA-DVTFSFTQYSTDEER 106
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                                                                                                                                                    5.4%; Score 94; DB 4; Length 501; 21.7%; Pred. No. 0.26; tive 35; Mismatches 115; Indels
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                                                                                                                                                          128;
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                                            US-09-410-551B-23
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LENGTH: 920
TYPE: PRT
ORGANISM: Bacillus stearothermophilus
S-09-463-402-6
Sequence 23, Application US/09410551B Patent No. 6503737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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Patent No. 6596510
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CURRENT APPLICATION NUMBER: US/09/463,402
CURRENT FILING DATE: 2000-01-31
PRIOR APPLICATION NUMBER: DE19732829.6
PRIOR FILING DATE: 1997-07-30
PRIOR APPLICATION NUMBER: PCT/EP98/04723
PRIOR FILING DATE: 1998-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lubitz, Werner
APPLICANT: Resch, Stephanie
TITLE OF INVENTION: Secretion of Carrier-bound
TITLE OF INVENTION: and into the Extracellular
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                                                                                                                                                                                                                                                                                                                                145 SKPDPVIPDPPIDPPPGTGKYTCPFAIWSLEEVYEPPTKNRPWPIYNAVELQPREFDVAL 204
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                                                                                                                                                                                                                                                                                                                                                                                                                  86
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                                                                                                                         AYQNITRGDFAQFVYRAVNINAVPEIVEVTAVNSTTVKVTF----NTQIADVDFTNF
                                                                                                                                                                                                                                                                                                                                                                       ETKFGVYDE--ITRLDAAVILARVLKLD-VDNAKDAGFTDVPKDRAKYVNALVEAGVLNG 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GLLGAELRILRKKP-----SDTAKPA-----APGGGRAAQLKLSSCPSGRQPASILDVRS 282
                                                                                                                                                                AFGNIER-----FIY-LKSINAYCSLSDIAAYHADGVIVGFWRDPSSGGAIPFDFTKF 313
                                                                                                                                                                                                           KIIANRYKLKADDVKLPFT--
                                                                                                                                                                                                                                               KDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATYLATDQAMRDQKYDIREGKKP---G 261
                                                                                                                                                                                                                                                                                                                                                                                                                -TQYSTDEERAFVRTELAALLASPLLIDAIDQLNPAYWT-----LLIAGGGSG 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ÓFKSYRK-----FVATTATAAMVASAVAÞVVSAASFTDVAPQYKDAIDFLVSTGATKGKT 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GKRPS---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5.2%; Score 91.5; D
21.5%; Pred. No. 1.3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                        ----GAYDPLTR-----VEM-----A 133
                                                                                                                                                                                                         ----DVNDTWAPYVKAL--YKYEVTKGKTPTSFG 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 920;
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: synthase fragment
US-09-410-551B-23
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TITLE OF INVENTION: POLYKETIDE SYNTHASE ENZYMES AND REC
TITLE OF INVENTION: CONSTRUCTS THEREFOR
FILE REFERENCE: 30062-20026.00
CURRENT APPLICATION NUMBER: US/09/410,551B
CURRENT FILING DATE: 1999-10-01
PRIOR APPLICATION NUMBER: US 60/139,650
PRIOR FILLING DATE: 1999-06-17
PRIOR PILLING DATE: 1999-06-17
PRIOR PILLING DATE: 1999-03-11
PRIOR APPLICATION NUMBER: US 60/123,810
PRIOR APPLICATION NUMBER: US 60/123,748
PRIOR FILING DATE: 1999-03-11
PRIOR PILLING DATE: 1998-10-02
NUMBER OF SEQ ID NOS: 72
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CURRENT APPLICATION NUMBER: US/09/940,316B
CURRENT FILING DATE: 2001-08-27
PRIOR APPLICATION NUMBER: 09/410,551
PRIOR APPLICATION NUMBER: US 60/139,650
PRIOR APPLICATION NUMBER: US 60/139,650
PRIOR FILING DATE: 1999-08-17
PRIOR APPLICATION NUMBER: US 60/123,810
PRIOR FILING DATE: 1999-03-11
PRIOR APPLICATION NUMBER: US 60/102,748
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
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                                                                                                                                                                                                   APPLICANT: KHOSLA, CHAITAN
APPLICANT: SANTI, DANIEL
APPLICANT: WU, KAI
TITLE OF INVENTION: POLYKETIDES ENCODING THE
TITLE OF INVENTION: GENE CLUSTER
FILE REFERENCE: 30062-20026.11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: KOSAN BIOSCIENCES, APPLICANT: REEVES, CHRISTOPH APPLICANT: CHU, DANIEL
                                                                                                                                                                                                                                                                                                                                                             APPLICANT: KOSAN BIOSCIENCES, I APPLICANT: REEVES, CHRISTOPHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       680 RPDAVI 685
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                                                                                                                                                                                                                                                                                                                                                                                                                                   23, Application US/09940316B
5. 6759536
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KPDPVI 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SSVVFAERMAECAAALREFVDWDLFTVLDDPAVVDRVDVVQPASWAMMVSLAAVWQAAGV 679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VTFSFTQYSTD---EERAFVRTELAALLASPLLIDAIDQLNPAYWTLLIA----GGGSGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             --TVSVSQPSRNRKNYKVQVKIQNPTACT--------ANGSCDPSVTRQAYAD 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LGNIGKDGKQTLVLNPRG-----VNPTNGVASLSQAGAVPALEKRV------
                                                                                                                                                                                                                                                                                                                                          CHU, DANIEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.2%; Score 91; DB ilarity 19.9%; Pred. No. 3.2; Conservative 35; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CHRISTOPHER
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NUMBER OF SEQ ID NOS: 72
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 23
LENGTH: 1509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Synthetic PKS OTHER INFORMATION: synthase fragment -09-940-316B-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                 TELEFAX: 619-455-5110
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 495 amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/455,559
FILING DATE: 31-MAY-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/003,144
FILING DATE: 12-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                     NAME: WETHERELL, JR. PH.D.,
REGISTRATION NUMBER: 31,678
REFERENCE DOCKET NUMBER: PD
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
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APPLICANT: HUYNH, THANH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WIMBER OF SEQUENCES: 27

CORRESPONDENCE ADDRESS:

ADDRESSEE: SPENGLEY HORN JUBAS & LUBITZ

STREET: 1880 CENTURY PARK EAST, FIFTH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: PatentI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: LOU ANGELES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         / Match 5.2%; Score 91;
Local Similarity 19.9%; Pred. No.
les 37; Conservative 35; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                    TELEPHONE:
TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LE OF INVENTION: GROWTH DIFFERENTIATION FACTOR-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              561 VDRVAVAQTLARRTHFAHRAVLLGDTVITTPPADRPDELVFVYSGQGTQHPAMGEQ-LAD 619
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CALIFORNIA
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                                      495 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                           E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                         619/455-5100
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Patent No. 6
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                                                                INFORMATION FOR SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSE: Fish & Richardson, P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
                         SEQUENCE CHARACTERISTICS:
LENGTH: 495 amino acid
                                                                                                REFERENCE/DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619/678-5070
                                                                                                                                                                                   FILING DATE: 12-JAN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lee, Se-Jin
APPLICANT: Huynh, Thanh
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mes 77; Conserv
                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM CONTROL OPERATING SYSTEM:
                                                                                                                                                    NAME: Lisa A. Haile, REGISTRATION NUMBER:
                                                                                                                                                                                                                   APPLICATION NUMBER: 08/4
FILING DATE: 31-MAY-1995
APPLICATION NUMBER: 08/0
                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
COPOLOGY:
                                                                                  ELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22 LNPRGVNPTNGVASLSQAGAVPA-----LEKRVTVSVSQPSR-----
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USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GTGKYTCPFAIWSLEEVYEPPTKN-----RPWPIYNAVELQPREF-----DVAL
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                 amino
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                                  495 amino acids
                                                                                  619/678-5099
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                                                                                                                                                                                                                                                                                                                                                             IBM Compatible
SYSTEM: Windows95
                                                                                                                                                                                                                                                                                                                                                                                                   Diskette
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                                                                ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GROWTH DIFFERENTIATION FACTOR-5
                                                                                                                                                                                                                        08/003,144
                                                                                                                                                                                                                                                          08/455,559
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; MOLECULE TYPE:
; FRAGMENT TYPE:
US-09-145-060-10
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Best Local Similarity
"" ches 77; Conserve
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                                           PCT-US94-00657-10
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PCT-US94-00657-10
                                                                                                                                                                                                                                                                                            COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/00657
FILING DATE: 1/12/94
CLASSIFICATION:
ATTORNEY/ACTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 10, Application PC/TUS9400657
GENERAL INFORMATION:
APPLICANT: SE-JIN LEE
APPLICANT: SE-JIN LEE
APPLICANT: HYNH, THANH
TITLE OF INVENTION: GROWTH DIFFERENTIATION FACTOR-5
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: SPENSLEY HORN JUBAS & LUBITZ
STREET: 1880 CENTURY PARK EAST, FIFTH FLOOR
CITY: LOS ANGELES
CITY: LOS ANGELES
  Query Match
                                                                                                                 TELEFAX: 619-455-5110
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 495 amino acids
                                                           TOPOLOGY: 15
MOLECULE TYPE:
                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:

NAME: WETHERELL, JR. PH.D., JOHN R.
REGISTRATION NUMBER: 31,678
REFERENCE/DOCKET NUMBER: FD3256 CIP OF PD2280
                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 619/455-5100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         161 GTGKYTCPFAIWSLEEVYEPPTKN-----RPWPIYNAVELQPREF------DVAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 227 GLLGAELRILRKKPL-----DVAKPAVPSSGRVAQLKLSSCPSGRQPAALLDVRSVPGLD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    59 ----NRKNYKVQVKIQNPTACTANGSCD------PSVTRQAYA-DVTFSFTQYSTDEER 106
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                                                                                              amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FLVFGRTKKRD----LFFNEIKARSGQD----DKTVYEYLFSQRRKRRAPLANRQGKRPS
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ilarity 22.0%;
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internal
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  5.1%;
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Score 90;
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DB
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Length 495;
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; NUMBER OF SEQ ID NOS: 469
; SOFTWARE: FastSEQ for Window
; SEQ ID NO 312
; LENGTH: 1569
; TYPE: PRT
; ORGANISM: Escherichia coli
US-09-711-164-312
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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PRIOR FILING DATE: 1999-11-9
NUMBER OF GEO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: ELITRA.008A
CURRENT APPLICATION NUMBER: US/09/711,164
CURRENT FILING DATE: 2000-11-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Zyskind, Judith
TITLE OF INVENTION: GENES ESSENTIAL FOR MICROBIAL PROLIFERATION AND ANTISENSE THERETY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                           y Match 5.1%; Score 90; DB 4; Length 1569;
Local Similarity 20.4%; Pred. No. 4.5;
hes 81; Conservative 32; Mismatches 143; Indels 1
                                                                                                                                                                1156 VNNNATGNFKIFVQDTGVSPQSDDAMTLVKTGGGDASFTLGNTGGFVDLGTYEYVLKSDG
                                                                                                                                                                                                                                               1108 GDVYLSAQTAAPATTNGIYNTLTTNELSGSGNF------YLHTNVAGSRGDQLV 1155
                                                                                                                                                                                                                                                                                                                              1060 TLTLGNGVDTTLADLTAASRGSVWLNSNNSCAGTSNC-----EYRVN-----SLLLND 1107
                                                                                 1216 NSNWNLTNDVKPNPDPIPNPKPDPKPDPKPDPNPKPDP----TPDPTPTPVPEKRITPST 1271
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                                      189 --IYNAVELQPREFDVALKDLL-------GNTKWRDWDSRLSYTTFRGCRGNGYID
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                                                                                                                                                                                                                                                                                     63 YKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEBRAFVRTBLAALLASPLL 122
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AAVLNMAATLPLVFDAELNSIRERLNIMKASPHNNNVW-----GATYNT----RNNVTTD
                                                                                                                                                                                                                                                                                                                                                                    TVTLGNIGKDGKQTLVLNPRG---VNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKN
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                                                                                                                     ---KPDPV---IPDPPIDPPPGTGKYTCPFAIWSLEEVYEPPTKNRPWP--
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APPLICANT: Hees, Jrgen
FITILE OF INVENTION: Tuberculosis Vaccine
FILE REFERENCE: 16862PUS
CURRENT APPLICATION NUMBER: US/09/485,717
CURRENT FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: EP 97114614.7
PRIOR FILING DATE: 1997-08-22
PRIOR FILING DATE: 1998-08-12
PRIOR FILING DATE: 1998-08-12
PRIOR FILING DATE: 1998-08-12
VINMER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LENGTH: 626
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
ORGANISM: TIPORMATION: Description of Artificial Sequence: recombinant
US-09-485-717-2
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US-09-485-717-2
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Search completed: January 4, 2005, 09:35:42
Job time : 27.8269 secs
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Patent No. 6673353
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 31; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1379 GYASWEHESGFYLDGVVKLNRFKSNVAG---KMSSGGA 1413
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                                                                                   253 DĖMAYSESQLIA 264
                                                                                                                                                                  138 KDGNEYIVVEKKKKSINQNNADIQVVNAISSLTYPGALVKANSELVENQPDVLPVKRDSL 197
                                                                                                                            104 EERAFVRTELAA 115
                                                                                                                                                                                                                                                                                                                                    31; Conservative
                                                                                                                                                                                                         50 TVSVSQPSRNRKNYKVQVKIQNPTACTANGSCDPSVTR-----QAYADVTFSFTQYSTD 103
                                                                                                                                                                                                                                                                                            14 KDGKQTLVL-----NPRGVNPTNGVASLSQAGA------VPALEKRV 49
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Result
No.
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Maximum
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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80
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1733.5
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

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11: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

12: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

13: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

14: /cgn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*

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US-10-622-064-4
US-10-622-124-5
US-10-622-087-5
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HILLIANTE STATES	Similarity 9; Conserva	pplication US2003015 ATION: Chmann, Mar chmann, Mar ennings, Ga ennings E: 1700.036 E: 1700.036 E: 1700.001 TION NUMBER DATE: 2002- TION NUMBER DATE: 2001- TION NUMBER DATE:		$\begin{array}{c} 55\\ 55\\ 55\\ 55\\ 55\\ 55\\ 55\\ 55\\ 55\\ 55$
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MAKLETVILGNIGKDGKQTLVLNPRGVNPINGVASLSQAGAVPALEKRVTVSVSQPSRNR 60	,	of Allergic Eosinophilic Diseases		Sequence 16, Appl Sequence 7, Appl Sequence 7, Appl Sequence 10, Appl Sequence 20, Appl Sequence 20, Appl Sequence 167, App Sequence 167, Appl Sequence 167, Appl Sequence 21, Appl Sequence 11, Appl Sequence 12, Appl Sequence 13, Appl Sequence 14, Appl Sequence 10, Appl Sequence 11, Appl Sequence 10, Appl Sequence 10, Appl Sequence 11, Appl Sequence 12, Appl Sequence 13, Appl Sequence 14, Appli Sequence 14, Appli Sequence 23, Appli Sequence 24, Appli Sequence 24, Appli

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                                                                                                                                                                                                                                            Query Match 100.0%; Score 1749; DB 14; Best Local Similarity 100.0%; Pred. No. 3.8e-161; Matches 329; Conservative 0; Mismatches 0;
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                181 PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY
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                                                           LLIDAIDQLNPAYWTLLIAGGGSGSKPDPVIPDPPIDPPPGTGKYTCPFAIWSLEEVYEP
                                                                                                                      KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASP
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NUMBER OF SEQ ID NOS: 350

SOFTWARE: PATENTIN Ver. 2.1

SEQ ID NO 217

LENGTH: 329

TYPE: PRT

ORGANISM: Bacteriophage Q-beta
US-10-050-898-217
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Publication No. US20030175711A1
GENERAL INFORMATION:
APPLICANT: Renner, Wolfgang A.
APPLICANT: Bachmann, Martin
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Best Local Similarity 100.
Matches 329; Conservative
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PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/331,045
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CURRENT APPLICATION NUMBER: US/10/050,898
CURRENT FILING DATE: 2002-01-18
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Sebbel, Peter
Piossek, Christine
Ortmann, Rainer
Luond, Rainer
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Tissot, Alain
Maurer, Patrick
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                                                             LATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP
                                                                                                                        PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY
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RESULT 4 US-10-346-190-11

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RESULT 5
US-10-465-811-2
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CURRENT FILING DATE: 2003-01-17
PRIOR APPLICATION NUMBER: 60/396,590
PRIOR FILING DATE: 2002-07-18
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR FILING DATE: 2002-07-08
PRIOR APPLICATION NUMBER: 60/389,898
PRIOR FILING DATE: 2002-06-20
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR FILING DATE: 2002-01-18
                                                                                                                   GENERAL INFORMATION:
APPLICANT: BACHMANN, MARTIN F
APPLICANT: RENNER, WOLFGANG I
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APPLICANT: Maurer, Patrick
APPLICANT: Pellicioli, Erica
APPLICANT: Renner, Wolfgang A.
TITLE OF INVENTION: Prion Protein Carrier-Conjugates
APPLICANT: RENNER, WOLFGANG A TITLE OF INVENTION: PACKAGED VIRUS-LIKE PARTICLES FOR USE AS ADJUVANTS: TITLE OF INVENTION: METHOD OF PREPARATION AND USE FILE REFERENCE: 1700.0290004

CURRENT APPLICATION NUMBER: US/10/465,811

CURRENT FILING DATE: 2003-06-20
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TYPE: PRT
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1 MAKLETYTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR

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; PRIOR APPLICATION NUMBER: US 60/389,898
; PRIOR FILING DATE: 2002-06-20
; NUMBER OF SEQ ID NOS: 131
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; SEQ ID NO 2
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Bacteriophage Q-beta
US-10-465-811-2
                                                                            ; LENGTH: 329
; TYPE: PRT
; ORGANISM: Bacteriophage Q-beta
US-10-289-456-11
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CURRENT APPLICATION NUMBER: US/10/289,456

CURRENT FILING DATE: 2002-11-07

PRIOR APPLICATION NUMBER: PCT/IB02/00166

PRIOR FILING DATE: 2002-01-21

PRIOR APPLICATION NUMBER: US 10/050,902

PRIOR FILING DATE: 2002-01-18

PRIOR PLING DATE: 2002-01-19

PRIOR APPLICATION NUMBER: US 60/396,635

PRIOR FILING DATE: 2002-07-19

PRIOR FILING DATE: 2001-11-07

NUMBER OF SEQ ID NOS: 170

SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 11, Application US/10289456 Publication No. US20040033211A1 GENERAL INFORMATION:
                                                                                                                                                          SEQ ID NO 11
  Matches
                                    Query Match
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APPLICANT: Maurer, Patrick
APPLICANT: Spohn, Gunther
TITLE OF INVENTION: Antigen Arrays for Treatment of Bone Disease
                    Local
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                h 100.0%; Score 1749;
Similarity 100.0%; Pred. No. 3.
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GENERAL INFORMATION:

APPLICANT: Bachmann, Martin F

APPLICANT: Maurer, Patrik F

ITITLE OF INVENTION: Hapten-Carrier Conjugates and Uses Thereof

FILE REFERENCE: 1700.0300001

CURRENT APPLICATION NUMBER: US/10/622,064

CURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: US 60/396,575

PRIOR RILING DATE: 2002-07-18

NUMBER OF SEQ ID NOS: 33

SOFTWARE: Patentin version 3.2

SEQ ID NO 4

LENGTH: 329

TYPE: PRT
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Publication No. US20040059094A1
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Tocal Similarity
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                  SSGGAIPFDFTKFDKTKCPIQAVIVVPRA 329
SSGGAIPFDFTKFDKTKCPIQAVIVVPRA 329
                                                              LATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP
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APPLICANT: Betchmann, Martin F
APPLICANT: Betchmann, Martin F
APPLICANT: Fulurija, Alma
TITLE OF INVENTION: Ghrelin-Carrier Conjugates
FILE REFERENCE: 1700.0340001
CURRENT APPLICATION NUMBER: US/10/622,124
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,638
PRIOR FILING DATE: 2002-07-19
NUMBER OF SEQ ID NOS: 146
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LENGTH: 329
TYPE: PRI
ORGANISM: Bacteriophage Q-beta
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Matches 329;
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301
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                                                                     LATDQAVRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP
                                                                                            LATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP 300
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SSGGAIPFDFTKFDKTKCPIQAVIVVPRA 329
                   SSGGAIPFDFTKFDKTKCPIQAVIVVPRA 329
                                                                                                                                              PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY
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US-10-622-087-5
; Sequence 5, Application US/10622087
; Publication No. US20040141984A1
; GENERAL INFORMATION:
 APPLICANT: INFORMATION:
 APPLICANT: Tissot, Alain
 APPLICANT: Tissot, Alain
 APPLICANT: Ortmann, Rather
 APPLICANT: Staufenbiel, Matthias
 APPLICANT: Staufenbiel, Matthias
 APPLICANT: Frey, Peter
 TITLE OF INVENTION: Amyloid Beta 1-6 Antigen Arrays
 FILE REFERENCE: 1700.0350002
 CURRENT FILING DATE: 2003-07-18
 CURRENT FILING DATE: 2003-07-18
 PRIOR APPLICATION NUMBER: US 60/396,639
 PRIOR APPLICATION NUMBER: US 60/396,639
 PRIOR APPLICATION NUMBER: US 60/470,432
 PRIOR FILING DATE: 2003-05-15
 NUMBER OF SEQ ID NOS: 93
 SOFTWARE: PatentIn version 3.2
 SEQ ID NO 5
 LENGTH: 329

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                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles FILE REFERENCE: 1700.0210001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER: OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 11
LENGTH: 328
TYPE: PRT
ROANISM: Bacteriophage Q-beta
US-10-243-739-11
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US-10-243-739-11
; Sequence 11, Application US/10243739
; Publication No. US20030091593A1
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; ORGANISM: Bacteriophage Q-beta
US-10-622-087-5
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APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Lechner, Franziska
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                                                                                                                                                                                                                           Matches 328;
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                     LLIDAIDQLNPAYWTLLIAGGGSGSKPDPVIPDPPIDPPGTGKYTCPFAIWSLEEVYEP 180
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PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY 240

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APPLICANT: Cielens, India:
APPLICANT: Cielens, India:
APPLICANT: Renhofa, Regina
TITLE OF INVENTION: Packaging of Immunostimulatory Sub
TITLE OF INVENTION: Method of Preparation and Use
FILE REPERENCE: 1700.0220001
CURRENT APPLICATION NUMBER: US/10/244,065
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/318,994
PRIOR FILING DATE: 2001-09-14
NUMBER: OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 11
LENGTH: 328
TYPE: PAT
ORGANISM: Bacteriophage Q-beta
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Lipowsky, Gerad
Pumpens, Paul
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                                                           LATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHADGVIVGFWRDP
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                                                                                                     PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY
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Tissot, Alain
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99.7%; Pred. No. 1.2e-159;
tive 0; Mismatches 0;
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US-10-243-739-16
Sequence 16, Application US/10243739
Publication No. US20030091593A1
GENERAL INFORMATION:
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US-10-244-065-16
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SEQ ID NO 16
SEG ID NO 16
SENGTH: 329
TYPE: PRT
ORGANISM: Bacteriophage SP
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Lechner, Franziska
TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of
TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
FILE REFERENCE: 1700.0210001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT APPLICATION NUMBER: US/10/243,739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
                                                                                                                                                                                                                                                    APPLICANT: Bachmann, Martin F. APPLICANT: Storni, Tazio APPLICANT: Maurer, Patrick APPLICANT: Tissot, Alain APPLICANT: Schwarz, Katrin
                 APPLICANT: Meijerink, Edwin
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Pumpens, Paul
APPLICANT: Cielens, Indulis
APPLICANT: Renhofa, Regina
TITLE OF INVENTION: Packaging of Immunostimulatory Substances
TITLE OF INVENTION: Method of Preparation and Use
FILE REFERENCE: 1700.0220001
CURRENT PILING DATE: 2002-09-16
CURRENT FILING DATE: 2002-09-16
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 NFKVQIKLQNPTACTRD-ACDPSVTRSAFADVTLSFTSYSTDEERALIRTELAALLADPL 119
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No. US20030099668A1
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58.3%; Pred. No. 3
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3.4e-86;
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; PRIOR FILING DATE: 2002-04-22
PRIOR APPLICATION NUMBER: 60/318,994
PRIOR FILING DATE: 2001-09-14
; NUMBER OF SEQ ID NOS: 73
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 16
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Bacteriophage SP
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US-10-289-454-16
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                                                                            ; ORGANISM: Bacteriophage SP
US-10-289-454-16
Query Match
Best Local Sim
Matches 196;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16, Application US/102 Publication No. US20030157479A1 GENERAL INFORMATION:
                                                                                                                                      SOFTWARE: PatentIn
SEQ ID NO 16
LENGTH: 329
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                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
                                                                                                                                                                                                PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bachmann, Martin
                                                                                                                 TYPE: PRT
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                    Similarity
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                  58.3%;
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Score 978; DB 14;
Pred. No. 3.4e-86;
5; Mismatches 77;
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        9
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N

Conservative

45;

18;

Gaps

AKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 61

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RESULT 15

US-10-346-190-16

US-10-346-190-16

Sequence 16, Application US/10346190

Publication No. US20030219459A1
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; ORGANISM: RNA-phage SP Al protein
US-10-346-190-16
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PRIOR FILING DATE: 2002-07-08
PRIOR PPLICATION NUMBER: 60/389,898
PRIOR FILING DATE: 2002-06-20
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR PILING DATE: 2002-01-21
PRIOR PPLICATION NUMBER: 10/050,902
PRIOR FILING DATE: 2002-01-18
NUMBER OF SEQ ID NOS: 164
                                                                                                                                                                                                                                                                          Query Match 55.9%; Score 978; DB 14; Best Local Similarity 58.3%; Pred. No. 3.4e-86; Matches 196; Conservative 45; Mismatches 77;
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SEQ ID NO 16
LENGTH: 329
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CURRENT APPLICATION NUMBER: US/10/346,190
CURRENT FILING DATE: 2003-01-17
PRIOR APPLICATION NUMBER: 60/396,590
PRIOR FILING DATE: 2002-07-18
PRIOR FILING DATE: 2002-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Maurer, Patrick
APPLICANT: Pellicioli, Erica
APPLICANT: Renner, Wolfgang A.
TITLE OF INVENTION: Prion Protein Carrier-Conjugates
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APPLICANT: Maurer, Patrick
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                                                                                122 LIDAIDQLNPAYW-TLLIAGGGSGSKPD----PVIPDPPIDPPGTGKYTCPFAIWSLEE 176
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NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL 121
                                                                                                                                                                                                    AKLNQVILSKIGKNGDQTLTLTFRGVNPINGVASLSEAGAVPALEKRVIVSVAQPSRNRX 60
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                                                                                                                               NEKVOIKLONPTACTRD-ACDPSVTRSAFADVTLSFTSYSTDEERALIRTELAALLADPL 119
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                                                                                                                                                                                                                                                                                                              Length 329;
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Job time: 82.5897 secs
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234 IDLDATYLATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSI--NAYCSLSDIAAYHADG 291
                                                                        292 VIVGFWRDPSSGGAIPFDFTKFDKTKCPIQAVIVVP 327
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MVIGEWID-SKSPQLPIDFTQFNSANCPVQTVIIIP 328
                                                                                                                                                                 IDLDATAMQSDDFVLSGRYGVRKVKFPGAFGSIK---YLLNIQGDAWLDLSEVTAYRSYG
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Title:
Perfect score:
                                                                             Database
                                                                                                             Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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PIR_79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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1749
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

3

SUMMARIES

Result No.

C;Date: 29-Jul-1981 #sequence_revision 24-Sep-1981 #text_change 09-Jul-2004	listeriolysin O pr	1 529 2 AC1100	89.5 5.	
ge Q	listeriolysin O pr	1 529 2 A43505	89.5	
coat protein - phage Q-beta	listeriolysin prec	N	89.5	
VCBPQB	probable tellurium	N	89.5	
RESULT 2	dutt1 protein - mo	1 1612 2 T30805	90 5.	
	hypothetical prote	N	90 5.	
	bone morphogenetic	1 495 2 S43294	90 5.	
Db 294 GMVIGFWTD-SKSPQLPTDFTQFNSANCPVQTVIIIP 329	major merozoite su	N	90.5 5.	
	hypothetical prote	N	90.5 5.	
Qy 291 GVIVGFWRDPSSGGAIPFDFTKFDKTKCPIQAVIVVP 327	hypothetical prote	2 836 2 C85858	90.5 5.	
	hypothetical prote	2 836 2 A98014	90.5 5.	
Db 237 YIDLDATAMQSDDFVLSGRYGVRKVKFPGAFGSIKYLLNIQGDAWLDLSEVTAYRSY 2	transcription fact	2 280 2 A53027		
	peptidase, M23/M37	2 609 2 G87496		
Qy 233 YIDLDATYLATDQAMRDQKYDIREGKKPGAFGNIERFIYLKSINAYCSLSDIAAYHAD 2	hypothetical prote	2 1461 2 A85547	91.5 5.	
	hypothetical prote	N	91.5	
Db 178 SIYEVGKEGSP-DIYERGDEVSVTFDYALEDFLGNTNWRNWDQRLSDYDIANRRRCRGNG 2	hypothetical prote	N	92 5.	
<u></u>	polymorphic membra	N	92.5 5.	
Qy 176 EVYEPPTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNG 23	optic lobe develop	3 974 1 A40213	93 5.	
	glutamine syntheta	N	93	
Db 120 LIVDAIDNLNPAYWAALLVASSGGGDNPSDPDVPVVPDVKPPDGTGRYKCPFACYRLG 1	hypothetical prote	N	93.5	
	growth/differentia	N	94 5.	
Qy 121 LLIDAIDQLNPAYW-TLLIAGGGSGSKPDPVIPDPPIDPPPGTGKYTCPFAIWSLE :	seeligeriolysin -	N	94.5 5.	
	cellulase (EC 3.2.	4 915 2 A43802	95 5.	
~	probable ATP-depen	N	95.5 5.	
	probable secreted	1 973 2 T35238	106 6.	
Qy 61 KNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASP 1	conserved hypothet	5 409 2 D90268	113.5 6.	
	coat protein - pha	7 131 1 VCBPP1	117 6.	
Db 1 MÁKLNQVTLSKIGKNGDQTLTLTPRGVNPTNGVÁSLSEÁGÁVPÁLEKRVTVSVAQPSRNR 6	coat protein - pha	9 132 1 VCBPQB	663 37.	
	readthrough protei	2 331 2 S01964	983 56.2	
Qy 1 MAKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNR 6		:		
	Description	Match Length DB ID	Score Matc	
Matches 197: Conservative 45: Mismatches 77: Indels 18: Gaps		≺	Query	
Best Local Similarity 58.5%; Pred. No. 4.8e-75;			di	

119

177

236 232

Query Match Best Local Similarity 58.5%; Pred. No. 4.8 Matches 197; Conservative 45; Mismatches Qy 1 MAKLETVTLGNIGKDGKQTLVLNPRGVNPTNGV	S01964 S01964 Creathrough protein - phage SP Cryspecies: S01964 Rrinokuchi, Y.; Jacobson, A.B.; Hirose, T.; Inayama, S.; Hirose, T.; Inayama, S.; Hirose, Title: Analysis of the complete nucleotide sequence of the A; Action Solids of the complete nucleotide sequence of the A; Accession: S01964 A; Accession: S01964 A; Status: preliminary A; Molecule type: genomic RNA A; Residues: 1-331 < INO> A; Cross-references: UNIPROT: P09677; EMBL: X07489 A; Note: the authors translated the readthrough stopcodon TGA C; Superfamily: phage GA coat protein	30 89 5.1 949 2 T08658 31 89 5.1 1039 2 S02711 32 88.5 5.1 1307 2 T35944 33 88.5 5.1 1307 2 T35944 34 88 5.0 928 2 D81411 36 87 5.0 346 2 B60340 37 87 5.0 346 2 H70567 38 86.5 4.9 4.9 29 2 B44997 39 86.5 4.9 593 2 F64523 40 86 4.9 130 1 A46324 41 86 4.9 130 1 A46324 42 86 4.9 933 2 T41703 43 86 4.9 933 2 S41539 45 85.5 4.9 312 2 A56911 ALIGNMENTS
56.2%; Score 983; DB 2; Length 331; similarity 58.5%; Pred. No. 4.8e-75; conservative 45; Mismatches 77; Indels 18; Gaps 9; MAKLETVILGNIGKDGKQTLVLNPRGVNPINGVASLSQAGAVPALEKRVTVSVSQPSRNR 60	#sequence_revision 19-Mar-1997 #text_change 09-Jul-2004 bson, A.B.; Hirose, T.; Inayama, S.; Hirashima, A. 6, 6205-6221, 1988 the complete nucleotide sequence of the group IV RNA coliphage SP. \$01963; MUID:88289362; PMID:3399390 YOUNIC RNA NO- UNIPROT:P09677; EMBL:X07489 translated the readthrough stopcodon TGA for residue 133 as Trp GA coat protein	hypothetical prote cellulase (EC 3.2. MKT1 protein - yea probable beta-gala probable lipoprote probable lipoprote transposase - Myco transposase - Myco merozoite surface hypothetical prote coat protein - pha probable chain len dipeptidyl aminope gastric mucin MUC5 fibringen-binding TRADD protein - hu

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C;Accession: A04225

R;Dhasse, P; Vandekerckhove, J.S.; Van Montagu, M.;
Eur. J. Biochem. 94, 375-386, 1979

A;Title: The primary structure of the coat protein A;Reference number: A04225; MUID:79148387; PMID:107

A;Accession: A04225

A;Molecule type: protein
A;Residues: 1-131 <DHA>
A;Cross-references: UNIPROT:P03616
C;Superfamily: phage GA coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Accession: A92221
A;Molecule type: protein
A;Residues: 1-60 <3TO>
A;Residues: 1-60 <3TO>
R;Maita, T.; Konigsberg, W.
1. Biol. Chem. 246, 5003-5024, 1971
A;Title: The amino acid sequence of the Obeta coat protein.
A;Reference number: A92088; MUID:71288580; PMID:5570434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:P03615
R;Stoll, E.; Wilson, K.J.; Reiser, J.; Weissmann, C.
J. Biol. Chem. 252, 990-993, 1977
A;Title: Revised amino acid sequence of Qbeta coat protein
A;Reference number: A92221; MUID:77118576; PMID:838709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: A92240; A92221; A92088; A04224
R;Bscarmis, C.; Sastry, P.A.; Billeter, M.A.
J. Biol. Chem. 253, 8390-8399, 1978
A;Title: Determination of the first half of the coat protein cistron of bacteriophage A;Reference number: A92240; MUID:79048469; PMID:361741
A;Accession: A92240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               coat protein - phage PRR1
c/Species: phage PRR1
c/Species: phage PRR1
c/Date: 30-Nov-1979 #sequence_revision 30-Nov-1979 #text_change 09-Jul-2004
c/Accession: A04225
R/Dhaese, P./ Vandekerckhove, J.S.; Van Montagu, M.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Accession: A92088
A;Molecule type: protein
A;Mesidues: 1-21,'D',23-55,57-132 <MAI>
C;Superfamily: phage GA coat protein
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A; Residues: 1-80 < ESC>
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VCBPP1
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Best Local S
Matches 132
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Best Local
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                                           124 DAIDQLNPAY 133
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                                                                                                                              68 KIQNPT--ACTANGSCDPSVTRQAYADVTFSFTQYSTDEER-AFVRTELAALLASPLLI- 123
                                                                                                                                                                                                 4
                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LIDAIDQLNPAY 132
                                                                                     KLVVPVVQSQTVNGIVTPVVVRTSYVTVDFDYDARSTTKBRNNFVGMIADALKADLMLVH
                                                                                                                                                                            QNLVLKDREATPNDHTFVPRDIRDNVGEVVESTGVPIGESRFTISLRKTSNGR--YKSTL
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                                                                                                                                                                                                                                                               6.7%; Score 117; DB 1; Length 131; 32.3%; Pred. No. 0.011; ative 15; Mismatches 57; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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probable secreted cellulase - Streptomyces coelicolor (;Species: Streptomyces coelicolor (;Species: Streptomyces coelicolor (;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 09-Jul-2004 (;Accession: T35238 R;Seeger, K.J.; Harris, D.; Parkhill, J.; Barrell, B.G.; Rajandream, M.A. submitted to the EMBL Data Library, September 1998 A;Reference number: Z21572 A;Accession: T35238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conserved hypothetical protein [imported] - Sulfolobus solfataricus
C;Species: Sulfolobus solfataricus
C;Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change 09-Jul-2004
C;Accession: D90268
R;She, Q.; Singh, R.K.; Confalonieri, F.; Zivanovic, Y.; Allard, G.; Awayez, M.J.; Chan-V.
Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; Redder, P.
arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J.
submitted to GenBank, April 2001
A;Description: Sulfolobus solfataricus complete genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:Q97YZ7; GB:AE006641; NID:g13814341; PIDN:AAK41403.1; GSPDB:Gt
C;Genetics:
A;Gene: SSO1152
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D90268
                                                                                                                                                                           A;Molecule type: DNA
A;Residues: 1-973 <SES>
A;Cross-references: UNIPROT:086728;
A;Experimental source: strain A3(2)
C;Genetics:
A;Gene: SCOEDB:SC5C7.31c
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A; Accession: D90268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 6.9
Best Local Similarity 23.0
Matches 62; Conservative
                                                                                           Matches
                                                                                                              Query Match
Best Local
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                                            138 IAGGGSGSKPDPVIPDPPIDPPGTGKYTCPFAIWSLEEVYEPPTKN-----RPWPIYNA 192
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                                                                                                                 Similarity
IAGGATNSWAGR-YATPPAGTPTFYGMY-----YDEKPVYHDPPSNQWFGFQAWSMERV 769
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AY-----WTLLIAGGGSGSKPDPVIPDPPIDPP------PGT------GKYTCPF 169
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                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6.5%; Score 113.5; DB 23.0%; Pred. No. 0.096;
                                                                                                            6.1%; Score 106;
23.4%; Pred. No. 1.
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A;Status: ProcessionA
A;Molecule type: DNA
A;Residues: 1-1326 <STO>
A;Cross-references: UNIPROT:Q9HYU6; GB:AE004752; GB:AE004091; NID:g9949424; PIDN:AAG0668
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R;Stover; C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probable ATP-dependent helicase PA3297 C;Species: Pseudomonas aeruginosa C;Date: 15-Sep-2000 #secmence revision
A43802
cellulase (EC 3.2.1.4) / cellulose 1,4-beta-cellobiosidase
N;Alternate names: endo-1,4-beta-glucanase
C;Species: Caldocellum saccharolyticum
C;Date: 30-Jan-1993 #sequence_revision 30-Sep-1993 #text_ch
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Date: 15-Sep-2000 #sequence_revision 15-Sep-2000
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                                                                                                                                                                                                                                                   -RDQKYDIREGK-KPGAFGNIERFI----YLKSINAYCS--LSDIAAYHADGVIVG----
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                                                                                                                                                                               -FWRDPSSG 303
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#sequence_revision 30-Sep-1993 #text_change 15-Mar-2004
                                                                                                                                              683
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24.9%; Pred. No. 15;
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    Pseudomonas aeruginosa (strain

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A.; Larbig,
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                                                         Caldocellum
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K.; Lim,
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R;Saul, D.J.; Williams, L.C.; Grayling, R.A.; Chamley, L.W.; Love, Appl. Environ. Microbiol. 56, 3117-3124, 1990
A;Title: celB, a gene coding for a bifunctional cellulase from the A;Reference number: A43802; MUID:91136262; PMID:2126700
A;Accession: A43802
                                                                                                                                                                                                                                                                                                                                                                                                                                            seeligeriolysin - Listeria seeligeri
C;Species: Listeria seeligeri
C;Date: 22-Nov-1993 #sequence_revision 01-Dec-1995 #text_change 09-Jul-2004
C;Accession: S22340
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S22340
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                                                                                                                                                                                                                                                                                                                                                                        R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 80-84, 11992
A;Title: Listeriolysin genes: complete sequence of ilo from Listeria ivanovii
A;Reference number: S22340; MUID:92182018; PMID:1543752
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C; Function:
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A; Residues: 1-915 <SAU>
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                                                                                                                                                                                                                                                                                                                                        A;Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 h 5.4%; Score 95; DB 2;
Similarity 22.1%; Pred. No. 10;
                                                                   TVSVSQPSRNRKNYKVQVKIQNPTACTANGSCDPSVTR-----QAYADVTFSFTQYSTD 103
                                                                                                                                         KDGKQTLVL--NPRGVNPT-----NGVASLSQAGA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EPST 501
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 EERAFVRTELAA
                                                                                                          KDGSEYIVVEKKKKGINQNNADISVINAISSLTYPGALVKANRELVENQPNVLPVKRDSL
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                                      TLSVDLPGMTKKDNKIFVK--NPTKSNVNNAVNTLVERWNDKYSKAYPNINAKI-DYS--
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                                                                                                                                                                              Conservative
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                                                                                                                                                                                         25.4%;
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                                                                                                                                                                                               Score 94.5; DI
Pred. No. 5.4;
                                                                                                                                                                              Mismatches
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residue 287
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A;Cross-references: UC;Genetics: C;Genetics: ATSP:F316.9
A;Gene: ATSP:F316.9
A;Map position: 1
                                                                                                                     C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 01-Feb-1999 #sequence_revision 01-Feb-1999
C;Accession: T00646
R;Federspiel, N.A.; Palm, C.J.; Conway, A.B.; Kurl; Vysotskaia, V.S.; Yu, G.; Ecker, J.; Theologis, submitted to the EMBL Data Library, February 1998
A;Reference number: Z14197
A;Accession: T00646
A;Status: translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                            hypothetical protein F316.9 - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 01-Feb-1999 #sequence_revision 01-Feb-1999
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C;Species: Homo sapiens (man)
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A; Residues: 1-540 <FI
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Best Local S
Matches 77
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;Date: 20-Feb_1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
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                                                               UNIPROT: 048683; EMBL: AC002396; NID: g2749918; PIDN: AAC00577.1;
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Pred. No.
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                                                                                                                                                                                                       A.B.; Kurtz, D.B.; Conway, A.R.; Au, Theologis, A.; Davis, R.W.
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N,Alternate names: omb protein
C;Species: Drosophila melanogaster
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 09-Jul
C;Accession: A40213; S34827
R;Pflugfelder, G.O.; Roth, H.; Poeck, B.; Kerscher, S.; Schwarz, H.; J
Proc. Natl. Acad. Sci. U.S.A. 89, 1199-1203, 1992
A;Title: The lethal(1)optomotor-blind gene of Drosophila melanogaster
A;Reference number: A40213; MUID:92159016; PMID:1741374
A;Accession: A40213
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A36911
glutamine synthetase I - Pyrococcus woesei
C;Species: Pyrococcus woesei
C;Species: 07-Apr-1994 #sequence_revision 18-Nov-1994 #text_change
                               A;Note: sequence extracted from NCBI backbone R;Poeck, B.; Balles, J.; Pflugfelder, G.O. Mol. Gen. Genet. 238, 325-332, 1993
A; Title: Transcript A; Reference number:
                                                                                                            A;Status: preliminary; not compared with conceptual A;Molecule type: mRNA A;Residues: 1-974 <PFL>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-439 <TIB>
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A;Reference number: A36911; MUID:93259940; PMID:8098326
A;Accession: A36911
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R;Tiboni, O.; Camma
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                                                                         A;Cross-references: UNIPROT:Q24432; GB:M81796; NID:g158018; A;Note: sequence extracted from NCBI backbone (NCBIP:82056)
                                                                                                                                                                                                                                                                                                                                            optic lobe development omb protein -
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Best Local
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2961-2969, 1993
identification in the 334827; MUID: 93261414;
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Pred. No.
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Pred. No. 6.7;
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optomotor-blind; PMID:8492800
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A;Accession: S34827
A;Molecule type: DNA
A;Residues: 1-447 <POE>
A;Cross-references: GB:S61732; NID:g402317; PIDN:AAB26697.1; PID:g402318
A;Experimental source: larva
C;Genetics:
A;Gene: FlyBase:bi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hickey, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzberg, Nucleic Acids Res. 28, 1397-1406, 2000
A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.
A;Reference number: A81500; MUID:20150255; PMID:10684935
A;Accession: A81723
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: GB:AE002293; GB:AE002160; NID:g7190298; PIDN:AAF39133.1; PID:g719030; A;Experimental source: strain Nigg (MoPn)
C;Genetics:
A;Gene: TC0264
C;Superfamily: Polymorphic membrane protein H family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polymorphic membrane protein H family TC0264 [imported] - Chlamydia muridarum (strain Ni C;Species: Chlamydia muridarum, Chlamydia trachomatis MoPn C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 09-Dec-2002 C;Accession: A81723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: FlyBase:FBgn0000179
C;Superfamily: optic lobe development omb protein; T-box homology
C;Keywords: DNA binding
F;337-521/Domain: T-box homology <TBX>
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A; Residues: 1-983 <TET>
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mes 50; Conserv
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                                                                                           572 YDNPLFMNAVTIPFVTLVSLQTTGGVTTSAVTLNNADTAHYGYQGSWSADWRRPPLAPDP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         529 MSNRGSDSDK---LNPTHVSSSRAPLHLGHÄGRPPHLHPHAALLDNQQDDDDKLLDVVGP 585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              91 ADVTF----
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                                                                                                                                                                                                                                    60 RKNYKVQVKIQN----PTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAAL 116
                                                                                                                                                                                                                                                                                                                                                                                    67;
                                                                                                                                                                                                                                                                                                                                  1 MAKLETVTLGNI-GKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRN 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9 LGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 68
SGMTPLD-----KSNTLYVTW------RPSSNYGVYKLDPQR----RGELVPN 669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADADVERDGSDSSCSESVGGSTGGAFRPTSTGSPKEAVGAAAAAAAAAG-----LNP
                                          ----PIDPPPGTGKYTCPFAIWSLEEVYEPPTKNRPWPIYNAVELQPREFDVALKDLLGN 210
                                                                                                                                          LASPLLIDAIDQLNPAYWTLLIAGG-----
                                                                                                                                                                                          ----KLGIDLESFLVPTYETÄKLGADTTVA------VNNNPTLDLVMANETEM 571
                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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Pred. No.
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                                                                                                                                                                                                                                                                                        528
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RESULT 15 1590696 16 Aypothetical protein ECs0541 [imported] - Escherichia coli (str. C. Species: Escherichia coli C. Species: Bscherichia coli C. Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change C. Accession: E90696 C. R. Hayashi, T., Makino, K., Ohnishi, M., Kurokawa, K., Ishii, K. Gasawara, N., Yasunaga, T., Kuhara, S.; Shiba, T., Hattori, M.,	175 BEVYEPPTKNRPWPIXNAVELQPREFDVALKOLLGNT	QY 70 QNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAF\RTELAALLASPLLIDAIDQL	Query Match S.3*; Score 92; DB 2; Length 385; Best Local Similarity 22.2*; Pred. No. 5.7; Matches 67; Conservative 40; Mismatches 127; Indels Qy 10 GNIGKDGKQTLVIAPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRI	: 1-385 <sto> ferences: UNIPROT:Q8X9Q4; GB:AE005174; NID:g12513977 ferences: Strain O157:H7, substrain EDL933 n: B85658 preliminary type: DNA : 1-385 <st2> ferences: GB:AE005174; NID:g12514497; PIDN:AAG55726. ferences: GB:AE005174; Substrain EDL933 : rA; terA_2</st2></sto>	RESULT 14 A85607 A85607 A85607 hypothetical protein terA [imported] - Escherichia coli (strain C;Species: Escherichia coli C;Species: Bscherichia coli C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change C;Accession: A85607; B85658 R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, iller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, E.; Nature 409, 529-533, 2001 A;Title: Genome sequence of enterohemorrhagic Escherichia coli (A;Reference number: A85480; MUID:21074935; PMID:11206551 A;Accession: A85607 A;Status: preliminary A;Molecule type: DNA	Qy 211 TKWRDWDSRLSYTTFRGCRGNGYIDLDATYLATDQAMRDQKYDIREGKKPGAFGNI
li (st _chang _hii, K	KWRDWDSRLSYTTFRGCRG 230	QYSTDEERAFVRTELAALLASPLLIDAIDQL 129	ngth 385 Indels TVSVSQPS	5 1	li (strain O157:H7, substrain EDL93: ext_change 09-Ju1-2004 ; Glasner, J.D.; Rose, D.J.; Mayhew, lanta, E.; Potamousis, K.; Apodaca, shia coli O157:H7.	AWRDQKYDIREGKKPGAFGNI 266

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DNA Res. 8, 11-22, 2001
A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli 0157:H7 and genc A;Reference number: A99629; MUID:21156231; PMID:11258796
A;Recession: E90696
A;Status: preliminary
A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-1461 <-HAY>
A;Ross-references: UNIPROT:08XD19; GB:BA000007; PIDN:BAB33964.1; PID:g13359999; GSPDB:GA;Experimental source: strain 0157:H7, substrain RIMD 0509952
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                                                                                                                                                                                                                                                                                              233 ----YIDLDATYLATDQAMRDQKYDIRE------GKKPG-----AFGNIERFIY 271
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             972 GVTINPVATDDIINAAEAGVAQTISGQVTGAEDGDT-----VTITLGGNTYTATVG---- 1022
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                916 NYTTTTDASGNWSVTVPASAVTALGQANYTVT----AAVTSDIGNSATASHNVLVDSALP 971
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HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0019028; F:structural molecule activity; IEA.
InterPro; IPR002703; Levi_coat.
InterPro; IPR002703; Levi_coat.
InterPro; IPR002504; RNA rec mot.
Pfam; PP01819; Levi_coat; 1.
PROSITE; PS00030; RRM RNP 1; UNKNOWN 1.
SEQUENCE 329 AA; 36134 MW; E0F30715A8C197CB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                            "Recombinant RNA phage Q beta capsid particles synthesized and self-assembled in Escherichia coli.";
Gene 137.133-137(1993).
EMBL; AY099114; AAM33127.1; -.
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Ball J.J., Ellington A.D.;
Bacher J.M., Bull J.J., Ellington A.D.;
"Evolution of phage with chemically ambiguous
BMC Evol. Biol. 3:24-24(2003).
[2]
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Bacteriophage Q-beta.
Viruses; sBRNA positive-strand viruses, no DNA stage; Leviviridae;
Allolevivirus.
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Kozlovska T.M., Cielens I., Dreilinna
Ose V., Pumpens P.;
"Recombinant RNA phage Q beta capsid p
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     PTKNRPWPIYNAVELQPREFDVALKDLLGNTKWRDWDSRLSYTTFRGCRGNGYIDLDATY
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Q8x045 neurospora
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p36687 pyrococcus
p36487 gars corona
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OSE V., Pumpens P.;
OSE V., Pumpens P.;
"Recombinant RNA phage Q beta capsid particles
assembled in Escherichia coli.";
Gene 137:133-137(1993).
Gene 137:133-137(1993).
EMBL; M99039; AAA16663.1; -.
EMBL; M99039; AAA16663.1; -.
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MEDLINE=94109687; PubMed=7506687;
Kozlovska T.M., Cielens I., Dreil
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GO; GO:0005198; F:structural molecule activity; IEA.

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R InterPro; IPR000504; RNA_rec_mot.

R InterPro; IPR000504; RNA_rec_mot.

R Pfam; PF01819; Levi_coat; 1.

R PGSITE; PS00030; RRM RNP 1; UNKNOWN 1.

PROSITE; PS00030; RNM, RNP 1; UNKNOWN 1.
      064303;
01-AUG-1998 (TrEMBLrel: 07, Creat 01-AUG-1998 (TrEMBLrel. 07, Last 01-MAR-2004 (TrEMBLrel. 26, Last Al-protein. Bacteriophage M11. Viruses; sRNA positive-strand v. Allolevivirus. Allolevivirus. NCBI TaxID=74336;
[1]
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Submitted (APR-1998) to the
EMBL; AF059242; AAC14700.1;
HSSP; P03615; 1QBE.
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MEDLINE=96190948; PubMed=8609616;

Meekwilder J., Nieuwenhuizen R., Pc

Beekwilder y., Nieuwenhuizen for the

"Secondary structure model for the

Control of A-protein synthesis.";

J. Mol. Biol. 256:8-19(1996).
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MEDILINE=9539761; PubMed=7723040;

Beekwilder M.J., Nieuwenhuizen R.,

"Secondary structure model for the
"Secondary Structure model for the
"RNA phage Q beta.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LLIDAIDQLNPAYWTLLIAGGGSGSKEDPV---IPDPPIDPPPGTGKYTCPFAIWSLEEV
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Last sequence up
Last annotation
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Pred. No. 9.1e-102;
6; Mismatches 46;
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DT 01-MAR-19;
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DE Readthroug
OS Bacteriopi
OC Viruses;
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01-MAR-1989 (Rel. 10,
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Readthrough protein F
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PROSITE;
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HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi_coat.
InterPro; IPR002703; RNA rec_mot.
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SEQUENCE FROM N.A.
MEDLINE=88289362; PubMed=3399390;
Hirashima A., Hirose T., Inayama S.,
"Analysis of the complete nucleotide
                                                                                           Viruses; ssRNA positive-strand
                                                                                                         Bacteriophage SP.
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J. Mol. Biol.
                                                                                 Allolevivirus
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Pred. No. 3.1e-100;
1; Mismatches 54;
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01-MAR-2004
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064310;
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Nucleic Acids Res. 16:
-!- FUNCTION: Not yet
-!- MISCELLANEOUS: The
                                                      MEDLINE=95239761; PubMed=7723040; Beekwilder M.J., Nieuwenhuizen R., "Secondary structure model for the
                                                                                                                                                   NCBI_TaxID=75725;
                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                              Viruses; ssRNA
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                    A phage Q beta.";
Mol. Biol. 247:9
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MEDLINE-96190948; PubMed=8609616;
Beekwilder J., Nieuwenhuizen R., Pc
"Secondary structure model for the
"Secondary structure mynthesis.";
Takel Befa 126.0010/1006/
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Submitted (APR-1998) to the
EMBL; AF059243; AAC14704.1;
HSSP; P03615; 1QBE.
                                                                                                                                                                     EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                  Viruses; sBRNA positive-strand viruses, no Allolevivirus; Allolevivirus subgroup III. NCBI_TaxID=12009;
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                                                                                                                                                                                             Bacher J.M., Bull J.J., Ellington A. "Evolution of phage with chemically BMC Evol. Biol. 3:24-24(2003).
                                                                                                     SEQUENCE
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                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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GO:0005198; F:structural molecule activity; IEA.
erPro; IPR002703; Levi coat.
erPro; IPR000504; RNA_rec_mot.
im; PF01819; Levi coat; 1.
NSITE; PS00030; RRM NNP 1; UNKNOWN 1.
NSITE; PS00030; RRM NNP 1; ONES5F9408334410 CRC64;
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AY099114; AAM33126.1;
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he first three domains of Q beta RNA
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                                          This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content use by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
                                                                                                                             MEDLINE=96347354; PubMed=8736553; Golmohammadi R., Fridborg K., Bundule M., Valegard Golmohammadi R., Fridborg K., Bundule M., Valegard "The crysteal structure of bacteriophage Q-beta at 3 Structure 4:543-554(1996).
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MEDLINE=79048469; PubMed=361741;

MESCLARMIS C., Sastry P.A., Billeter M.A.;

"Determination of the first half of the
                                                                                                                                                                                                                                                                                                                                                                                                                                             "Recombinant RNA phage Q-beta capsid assembled in Escherichia coli.";
Gene 137:133-137(1993).
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21-JUL-1986
                     entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                Maita T., Konigsberg W.;
"The amino acid sequence of the Q-beta
J. Biol. Chem. 246:5003-5024(1971).
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Stoll E., Wilson K.J., Reiser J.
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MEDLINE=96190948; PubMed=8609616;

MEDLINE=96190948; PubMed=8609616;

Beekwilder J., Nieuwenhuizen R., Poot R., van Duin J.;

"Secondary structure model for the first three domains
"Secondary structure model for the first three domains
"Control of A-protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3D-structure;
INIT_MET
CONFLICT :
                                                                             MEDLINE=95239761; PubMed=7723040; Beekwilder M.J., Nieuwenhuizen R., "Secondary structure model for the RNA phage Q beta."; J., Mol. Biol. 247:903-917(1995).
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01-JUN-2003 (TrEMBLrel.
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01-MAY-2000
01-MAY-2000
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MEDINE-96190948; PubMed-8609616;
Beekwilder J., Nieuwenhuizen R., Pc
"Secondary structure model for the
"Secondary structure mynthesis.";
Control of A-protein synthesis.";
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01-MAY-2000 (TrEMBLrel.
01-JUN-2003 (TrEMBLrel.
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HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
                                                                                                                                                                                                        GO; GO:0019028; C:viral capsid; IEA. GO; GO:0005198; F:structural molecule InterPro; IPR002703; Levi coat.
                                                                                                                                                                                                                                                                     Beekwilder J., Nieuwenhuizen R., Poot R., van
Submitted (MAR-1998) to the EMBL/GenBank/DDBJ
EMBL; AF052431, AAC06250.1; -
HSSP; P03615; 1QBE.
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                                                                                                                                                                       Pfam; PF01819; Levi_coat; Coat protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RNA
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(APR-1998) to the
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Pred. No. 3.7e
12; Mismatches
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Pred. No. 8.8e
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hree domains
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databases.
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JE Major coat protein.

Enterobacteria phage NL95.

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Best Local &
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P09673;
01-MAR-1989
01-MAR-1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Analysis of the complete coliphage SP.";
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Hirashima A., Hirose T., Inayama S., Inokuchi
"Analysis of the complete nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Coat protein.
Bacteriophage SP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic Acids Res. 16:6205-6221(1988).
-!- FUNCTION: Forms the phage shell; binds to the phage RNA.
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Best Local S
Matches 101
MEDLINE-9825058; PubMed-9589071;

X MEDLINE-9825058; PubMed-9589071;

A Ikeno S., Tsuji T., Higashide K., Kinoshita N., Hamada M., Hc
Ikeno S., Tsuji T., Higashide K., Kinoshita N., Hamada M., Hc
T "A 7.6kb DNA region from Streptomyces kasugaensis M338-M1 inc
some genes responsible for kasugamycin biosynthesis.";

J. Antibiot. 51:341-352(1998).

R EMBL; AB005901; BAA25656.1; -.

R EMBL; AB005901; BAA25656.1; -.

R HSSP; P27830; IBXK.

R GO; GO:0003824; F:catalytic activity; IEA.

R GO; GO:0005875; F:racemase and epimerase activity, acting on
GO; GO:0005975; P:carbohydrate metabolism; IEA.

R GO; GO:0005975; P:nucleotide-sugar metabolism; IEA.
                                                                                                                                                                                                                                                                                                      01-AUG-1998
01-AUG-1998
01-OCT-2003
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF059243; AAC14703.1; -.
HSSP; P036.15; 10BE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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Beekwilder J., Nieuwenhuizen
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J. Mol. Biol. 247:9
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NCBI_TaxID=75725;
                                                                                                                                                                                                                           Streptomyces kasugaensis.
Bacteria; Actinobacteria; Actinobacteridae; Actir
Streptomycineae; Streptomycetaceae; Streptomyces.
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STRAIN=M338-M1;
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GO:0005198; F:structural molecule
erPro; IPR002703; Levi_coat.
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Nieuwenhuizen R.,
ure model for the
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Pred. No. 4.4e-33;
0; Mismatches 21;
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ne first three
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last two domains
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Best Local
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COAT_BPPRR
P03616;
                                                                                                                                                                                                                                                     Pfam; PF01819; Levi coat; 1.
Coat protein; Direct protein and SEQUENCE 131 And Coat; 1.
                                                                                                                                                                                                                                                                                                         Eur. J. Blochem. 94:375-386(1979).
-I- FUNCTION: Forms the phage shell; binds to PIR; A04225; VCBEP1.
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Viruses; ssRNA positive-strand viruses,
unclassified Leviviridae.
NCBI_TaxID=12024;
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21-JUL-1986 (Rel.
05-JUL-2004 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR001509; Epimerase_Dh.
InterPro; IPR008099; Nuc sugar_epim.
Pfam; PF01370; Epimerase; 1.
PRINTS; PR01713; NUCEPIMERASE.
  Q97YZ7
                                                                                                                                                                                                                                                                                                                                                 bacteriophage PRR1."
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                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=79148387; PubMed=107028;
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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                                                                                                                                  KIQNPT--ACTANGSCDPSVTRQAYADVTFSFTQYSTDEER-AFVRTELAALLASPLLI- 123
                                                                                                                                                                                    QTLVLNPRGVNPT-----NGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQV
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                                                             DTIVNLOGVY 131
                                                                                                             KLVVPVVQSQTVNGIVTPVVVRTSYVTVDFDYDARSTTKERNNFVGMIADALKADLMLVH
                                                                                                                                                            QNLVLKDREATPNDHTFVPRDIRDNVGEVVESTGVPIGESRFTISLRKTSNGR--YKSTL 61
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14535 MW; E7E639E1E50FC612 CRC64;
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C;Accession: S22340
K;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete se
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A;Title: The primary structure of the coat protein of A;Reference number: A04225; MUID:79148387; PMID:10702
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Note: the authors translated the readthrough stopcodon TGA for residue
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                                                                                                                                                                                                                                                             DAIDQLNPAY
                                                                                                                                                                                                                                                                                                    KLVVÞVVQSQTVNGIVTÞVVVRTSYVTVDFDYDARSTTKERNNFVGMIADALKADLMLVH 121
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MUID:92182018; PMID:1543752
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Pred. No. 6.4e-45;
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C; Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_cnange C; Accession: 824231
R; Rasmussen, O.F.; Beck, T.; Olsen, J.E.; Dons, L.; Rossen, L. Infect. Immun. 59, 3945-3951, 1991
A; Title: Listeria monocytogenes isolates can be classified into A; Reference number: 824230; MUID:92040062; PMID:1937753
A; Accession: 824231
A; Status: nucleic acid sequence not shown; translation not shown A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Hence 1
A; Map position: 1
A; Introns: 14/3; 281/3;
C.Sumerfamily: pyruvate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Federspiel, N.A.; Palm, C.J.; Conway, A.B.; Kurtz, D.B.; Conway, Yysotskaia, V.S.; Yu, G.; Ecker, J.; Theologis, A.; Davis, R.W. submitted to the EMBL Data Library, February 1998
A;Reference number: Z14197
A;Accession: T00646
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypothetical protein F316.9 - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 01-Feb-1999 #sequence_revision 01-Feb-1999
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A;Molecule type: DNA
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A;Variety: strain 12067
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A; Residues: 1-540 < FED>
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                                                                                                                                                                                                                                                                                                               KPAAPMSKSPQGFATPRVYKPAPTKTSLSTSHSSLKKEK--
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dehydrogenase
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25.8%;
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%; Pred. No. 0.29
14; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23;
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Pred. No. 0.
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A;Residues: 1-529 <RAS>
A;Cross-references: UNIPROT:P13128; EMBL:X60035; NID:g44110; PIDN:CAA42639.1; PIDA;Experimental source: strain 12067, serotype 4b
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1991
                                                                                                                                                                                                                                                                               A;Status: preliminary
A;Molecule type: DNA
A;Residues: 413-480 <ME2>
A;Cross references: GB:M39171
R;Michel, E.; Reich, K.A.; Favier, R.; Berche, P.; Cossart,
Mol. Microbiol. 4, 2167-2178, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Listeria monocytogenes
C;Species: Listeria monocytogenes
C;Date: 21-Oct-1992 #sequence revision 21-Oct-1992 #text change 09-Jul-2004
C;Accession: A43505; S03306; A47606; S12400; A61079
R;Mengaud, J.; Vicente, M.F.; Chenevert, J.; Pereira, J.M.; Geoffroy, C.; Gicquel-Sanzey
Infect. Immun. 56, 766-772, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
                                                                                                                                                                    A;Title: Attenuated mutants of the intracellular bacterium Listeria A;Reference number: S12400; MUID:91211627; PMID:1965218 A;Accession: S12400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Experimental source: strain EGD
A;Note: this sequence is derived from a weakly hemolytic strain,
R;Mengaud, J.; Chenevert, J.; Geoffroy, C.; Gaillard, J.L.; Cosss
Infect. Immun. 55, 3225-3227, 1987
A;Title: Identification of the structural gene encoding the SH-ac
A;Reference number: A47606; MUID:88057627; PMID:2824384
A;Accession: A47606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 δ
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A; Experimental source: C; Genetics:
                                                                                         A; Molecule type: DNA
A; Residues: 483-493 <MIC>
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A;Accession: A43505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Title: Expression in Escherichia coli and sequence analysis of the listeriolysin O det
A;Reference number: A43505; MUID:88153053; PMID:3126142
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Molecule type: DNA; Residues: 1-529 < MEN>; Residues: 1-529 < MEN>; Residues: 1-529 < MEN>; Residues: 1-529 < MEN>; Residues: 1-529 < MEN>; Residues: 1-529 < MEN>; Residues: 1-529 < MEN>; Residues: 1-529 < MENOT: P13128; GB: M24199; NID: g149652; PIDN: AAA03018.1; PID: g149653; Note: this sequence is derived from a strongly hemolytic strain, serotype 1/2c; Pomann, E.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakraborty, T.; Chakrab
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;Residues: 1-529 <DOM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EERAFVRTELAA 114
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Pred. No. 0.71
23; Mismatches
                                        serotype 1/2d
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coat protein - phage JP34
C;Species: phage JP34
A;Note: host Escherichia coli
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
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A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; A;Title: Comparative genomics of Listeria species.

A;Reference number: AB1077; MUID:21537279; PMID:11679669

A;Accession: AC1100
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A; Cross references: UNIFROT: P13128; GB:NC_003210; PIDN: CAD00729.1; PID: 916409567; GSPDB:C
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                      Virology
                                                                                        C; Accession: A46324
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                      M.R.; Hirashima, A.; Van Duin, 170, 238-242, 1989
Nucleotide sequence from the
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urget, O.; Entian, K.D.; |
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RESULT 11
T31660
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VCBPGA
                                                C;Accession: T31660
R;Bird, A.P.; Clark, V.; Jones, S.J.; Leitgeb, S.; submitted to the EMBL Data Library, December 1996
A;Reference number: Z21049
A;Accession: T31660
                                                                                                                                hypothetical protein COS41.6 - sea squirt (Ciona in C;Species: Ciona intestinalis C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: The complete nucleotide sequence of the group II RNA coliphage GA A;Reference number: A92000; MUID:86223910; PMID:3711059 A;Accession: A29178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: phage GA
A;Note: host Escherichia coli
C;Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change 09-Jul-2004
C;Accession: A29178
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C;Superfamily: phage GA coat proteil
C;Keywords: coat protein
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A;Accession: A46324
A;Molecule type: genomic RNA
A;Residues:
                A;Status: preliminary; translated A;Molecule type: DNA
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A; Residues: 1-130 < INO>
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ivanolysin precursor - Listeria ivanovii
(;Species: Listeria ivanovii
(;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text
(;Accession: $22341; $33683
R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete sequence of ilo f
A;Reference number: $22340; MUID:92182018; PMID:1543752
     R;Johnson, D.
submitted to the EMBL Data Library, February
                                                                                                    RGBYH1
CYC1/CYP3 transcription activator - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein L9672.1; protein YLR256w; regulatory prot
C;Species: Saccharomyces cerevisiae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Superfamily: dipeptide transport protein F;1-24/Domain: signal sequence #status predicted <SIG>F;25-528/Product: ivanolysin #status predicted <MAT>
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                                                                  C; Date: 30-Sep-1991;
C; Accession: S59400;
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A; Residues: 1-319, 'T', 321-528
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:P31831; EMBL:X60461
A;Note: the authors translated the codon ACA fo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: EMBL:X60461
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Accession: S36683
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               submitted to the EMBL Data Library, July 1991
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Best Local S
Matches 29
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number:
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                                                                                                                                                                                                                                                                 EERAFVRTELAA 114
                                                                                                                                                                                                                                                                                                    TLSIDLP--GMVNHDNEIVVQNATKSNINDGVNTLVDRWNNKYSEEYPNIS---AKIDYD
                                                                                                                                                                                                                                                                                                                                                                       KEGNOYIVVEKKKKSINQNNADIQVINSLASLTYPGALVKANSELVENQPDVLPVKRDSV
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                                                                                                                                                                                                                                                                                                                                                                                                          KDGKQTLVL-----NPRGVNPTNGVASLSQAGA----
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                                                                  #sequence_revision 23-Fe; A31312; S15447; S05804;
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28.3%;
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Pred. No. 8.
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                                                                                   23-Feb-1996 #text_change 12-Nov-1999
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                                                                    S15446
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                                                                                                                      regulatory protein CYP1;
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C;Superfamily: regulatory protein HAP1; GAL4 zinc binuclear cluster homology C;Keywords: DNA binding; heme binding; transcription regulation; zinc finger F;1-148/Domain: DNA binding #status predicted <DNA>F;59-98/Domain: GAL4 zinc binuclear cluster homology <GAL4>F;64-84/Region: zinc finger CCC motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Residues: 1-144,'I',146-322,'R',324-454,'N',456-507,'M',509-586,'K',588-882,'N',884-95
A;Cross-references: EMBL:J03152; NID:g171645; PIDN:AAA34662.1; PID:g171646
R;Creusot, F.; Verdiere, J.; Gaisne, M.; Slonimski, P.P.
J. Mol. Biol. 204, 263-276, 1988
A;Title: CYP1 (HAP1) regulator of oxygen-dependent gene expression in yeast. I. Overall A;Reference number: S15447; MUID:89125585; PMID:2851658
A;Accession: S15447
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A;Residues: 1-1502 <JOH>
A;Residues: 1-1502 <JOH>
A;Residues: 1-1502 <JOH>
A;Cross-references: EMBL:U20865; NID:g662330; PIDN:AAB67387.1; PID:g662331; GSPDB:GN0001
A;Experimental source: strain S288C (AB972)
A;Pfeifer, K.; Kim, K.S.; Kogan, S.; Guarente, L.
                                                                                                                                                                                                                          C;Date: 10-Oct-1995 #sequence revision 10-Nov-1995 #text_change 09-Jul-2004 C;Accession: S57337; S50279; S63024; S6307; S65096 R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M. submitted to the EMBL Data Library, June 1995
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A;Cross-references: SGD:S0004246; MI
A;Reference number: S57533
A;Accession: S57537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;177-189/Region: glutamine-rich
;245-445/Domain: heme binding #status predicted <HEM>
;299-304,323-328,347-352,373-378,389-394,415-420/Region: 6-residue repeats
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;Residues: 1-62,'R',64-1305,'Y',1306-1470,'LVDFYRADFPIWE';Cross-references: EMBL:X13793
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'Title: Functional dissection and sequence of yeast HAP1 activator.'Reference number: A31312; MUID:89106221; PMID:2643482
'Accession: A31312
                                                                                                                                                                                                                                                                                                                           Species: Saccharomyces cerevisiae
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                                                                                                                                                                                                                                                                                                                                                                               protein - yeast (Saccharomyces cerevisiae)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TLTSPQGSDLASNSATQYKPDLEDFLMQNSNFNGLMINPSSLVEVVGGYN 1447
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Pred. No. 28
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                                                                                                                                                                                                                                                                                                                                                     protein YNL085w
                                                                           EMBL:X89016; NID:g887621; J.D.; Wickner, R.B.
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                            satellite dsRNA of
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A;Molecule type: DNA
A;Residues: 1-29,'G', 31-808,'TMKTCLNYH' <VER>
A;Residues: 1-29,'G', 31-808,'TMKTCLNYH' <VER>
A;Cross-references: EMBL:U09129; NID:9520475; PIDN:AAC49470.1;
R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M.
submitted to the Protein Sequence Database, April 1996
                                                                                                                                                                                                                                                                                                   A;Title: Genome Sequence of the Plant Pathogen and Biotechnology A;Reference number: A97359; MUID:21608551; PMID:11743194 A;Accession: C98221
                                                                                                                                                                                                                                                                                                                                                                                  R;Goodner, B.; Hinkle, G.; Gattung, S.; Miller, N.; Blanchard, M.; Qurollo, B.; Goldi
A.; Liu, F.; Wollam, C.; Allinger, M.; Doughty, D.; Scott, C.; Lappas, C.; Markelz,
Science 294, 2323-2328, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein AGR_I_1428 [imported] - Agrobacterium tumefaciens (strain C58, Cerec C;Species: Agrobacterium tumefaciens C;Species: Agrobacterium tumefaciens C;Date: 22-Oct-2001 #text_change 09-Jul-2004 C;Date: 22-Oct-2001 #text_change 09-Jul-2004
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A;Title: The sequence of a 17 933 bp segment of Saccharomyces
A;Reference number: S65092; MUID:96310628; PMID:8740422
                                                                                                                                                                                                                    A; Molecule type: DNA
A; Residues: 1-136 < KUR>
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C;Superfamily: Saccharomyces cerevisiae MKT1 protein
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Yeast 12, 485-491, 1996
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A;Cross-references: EMBL:Z71361; MIPS:YNL085w
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A; Accession: S63017
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                                                                                                                                   A; Gene:
                                                                                                                                                                                          A; Cross-references:
                                                                                                                                                                                                                                                                            A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Accession: C98221
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11.7%; Score 77.5; I
25.4%; Pred. No. 2.2;
tive 14; Mismatches
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Copyright

GenCore version 5.1.6 (c) 1993 - 2005 Compugen

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Scoring table:

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                                                                                                                                                                                     MEDLINE=96347354; PubMed=8736553; Golmohammadi R., Fridborg K., Bundule M., Valegard K., Liljas L.; "The crystal structure of bacteriophage Q-beta at 3.5-A resolution."; Structure 4:543-554(1996).
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MEDILINE=77118576; PubMed=838709;

Stoll E., Wilson K.J., Reiser J., Weis
"Revised amino acid sequence of Qbeta
and 60.";
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"Recombinant RNA phage Q-beta capsid particles assembled in Bscherichia coli.";
Gene 137:133-137(1993).
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MEDLINE=94109687; PubMed=7506687;
Kozlovska T.M., Cielens I., Dreilinna
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                                                                                                                                                                                                                           Coat protein.
Bacteriophage Q-beta.
Viruses; ssRNA positive-strand
Allolevivirus; Allolevivirus su
NCBI_TaxID=12009;
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EMBL; V00643; CAA23992.1; -.
PIR; A92240; VCBPQB.
PDB; 1QBE; X-ray; A/B/C=1-13
                                                                                                    Bacher J.M., Bull J.J., Ellington A.D.;
"Evolution of phage with chemically ambiguous
BMC Evol. Biol. 3:24-24(2003).
EMBL; AY099114; AAM33126.1; -.
                                                                      SEQUENCE 13
                                                                                                                                                                              SEQUENCE FROM N.A.
PubMed=14667253;
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                  Similarity
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1819; Levi_coat; 1.
ure; Coat protein; Direct protein sequencing; RNA-binding.
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RESULT 4
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AC AAA1
DT 02-N

AAA16663; AAA16663; 02-MAR-2004

PRELIMINARY; (Trembirel.

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01-OCT-2002 (TrEMBLrel. 22, Last sequence
05-JUL-2004 (TrEMBLrel. 27, Last annotation of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of 
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PubMed=14667253;
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Bacher J.M., Bull J.J., Ellingt
"Eyolution of phage with chemic
"Eyolution of phage with chemic
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GO; GO:005198; F:structural molecule activity;
InterPro; IPR002703; Levi coat:
InterPro; IPR002504; RNA_rec_mot.
Pfam; PF01819; Levi_coat; 1.
PROSITE; PS000303; RRM_RNP 1; UNKNOWN 1.
SEQUENCE 329 AA; 36134 MW; E0F30215A8C197CB
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HSSP; P03615; 1QBE.
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Wozlovska T.M., Cielens I., Dreilinna
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                                                 LIDAIDQLNPAY 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 663; DB 2; ilarity 100.0%; Pred. No. 6.4e-56; Conservative 0; Mismatches 0;
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Best Local
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Beekwilder J., Nieuwenhuizen R., Poot R., van D
Beekwilder J., Nieuwenhuizen R., Poot R., van D
"Secondary structure model for the first three
Control of A-protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
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Gene 137:133-137(1993).
EMBL; M99039; AAA16663.1; -.
EMBL; M99039; AAA16663.1; -.
                                                                                                                             Beckwilder M.J., Nieuwenhuizen R., van Duin J.
Submitted (APR-1998) to the EMBL/GenBank/DDBJ
EMBL; AF059242; AAC14699.1; -
HSSP; P03615; 10BE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIINE=95239761; PubMed=7723040; Beekwilder M.J., Nieuwenhuizen R., "Secondary structure model for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q9TOR9;
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01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2003 (TrEMBLrel. 24,
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Viruses; ssRNA positive-strand
                    GO; GO:0019028; C:viral capsid; IEA. GO; GO:0005198; F:structural molecule InterPro; IPR002703; Levi_coat. Pfam; PF01819; Levi_coat; 1. Coat protein.
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Kozlovska T.M., Cielens I., Dreilinna D., Dislers A.,
Ose V., Pumpens P.;
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Best Local Similarity
Matches 111; Conserv
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GO; GO:005198; F:structural molecule activity;
InterPro; IPR002703; Levi_coat.
InterPro; IPR000504; RNA_rec_mot.
Pfam; PF01819; Levi_coat; 1.
PROSITE; P500030; RRM RNP 1; UNKNOWN 1.
SEQUENCE 329 AA; 35880 MW; 372B1DD5DDB52F15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beekwilder J., Nieuwenhuizen R., Po
"Secondary structure model for the
Control of A-protein synthesis.";
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Viruses; ssRNA positive-strand viruses,
Allolevivirus.
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HSSP; P03615; 1QBE.
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ d
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J. Mol. Biol. 247:903-917(1995).
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                                     LIDAIDQLNPAY 132
                                                                                      NYKVQVKIQNPTSCTASGTCDPSVTRSAYADVTPSFTQYSTDEERALVRTELKALLADPM
                                                                                                              NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL
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LIDAIDNLNPAY 133
                                                                                                                                                                                AKLQAITLSGIGKNGDVTLNLNPRGVNPTNGVAALSEAGAVPALEKRVTISVSQPSRNRK
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84.1%;
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tive 10; Mismatches 11
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Beekwilder J., Nieuwenhuizen R., Poot R.,
"Secondary structure model for the first t
Control of A protein synthesis.";
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Bacteriophage M11.
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GO:0005198; F:structural molecule
erPro; IPR002703; Levi_coat.
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HSSP; P03615; 1QBE.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; IEA.
InterPro; IPR002703; Levi_coat.
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
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[3]
                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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SEQUENCE
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HSSP; P03615; 1QBE.
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
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P09677;
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Pfam; PF01819; Levi_coat; 1.
Coat protein; RNa-binding.
SEQUENCE 132 AA; 14129 MW;
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01-MAR-1989 (Rel. 10, Last sequence update)
29-MAR-2004 (Rel. 43, Last annotation update)
Readthrough protein A1 [Contains: Coat protei
                                                                                                                                                                                                                                                                                                                                                                                                Nucleic Acids Res. 16:6205-6221(1988).
-I- FUNCTION: Not yet known.
-I- MISCELLANEOUS: The readthrough protein Al includes
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or send an email to license@isb-sib.ch).
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Pred. No. 3.4e-44;
2; Mismatches 14;
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Readthrough protein A1.
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Pred. No. 9.8
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Al-protein.
Enterobacteria phage NL95.
Viruses; ssRNA positive-strand viruses, no DNA stage; Leviviridae;
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"Secondary structure model for the
Control of A.protein synthesis.";
J. Mol. Biol. 256:8-19(1996).
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01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2003 (TrEMBLrel. 24,
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SEQUENCE 13
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GO; GO:0005198; P:structural molecule activity;
InterPro; IPR002703; Levi coat.
Pfam; PF01819; Levi coat; 1.
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HSSP; P03615; 1QBE.
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MEDLINE=95239761;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    h 75.6%; Score 501.5; DB 2; Length Similarity 75.8%; Pred. No. 1e-40;
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J. Mol. B:~"
                                                                                                                                                                                                                 COAT BPPRR
P03616;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=96190948; PubMed=8609616; Beekwilder J., Nieuwenhuizen R., Po "Secondary structure model for the Control of A-protein synthesis.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF01819; Levi coat; 1.
PROSITE; PS00030; RRM RNP 1; UNKNOWN
SEQUENCE 330 AA; 36175 MW; 961E5
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Submitted (APR-1998)
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                                                           Eur.
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InterPro; IPR000504; RNA_rec_mot.
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InterPro; IPR002703; Levi_coat. Pfam; PF01819; Levi_coat; 1. Coat protein; Direct protein se SEQUENCE 131 AA; 14535 MW;
                                          PIR;
                                                                     Dhaese P., Vandekerckhove J., "The primary structure of the bacteriophage PRR1.";
                                                                                                                                 NCBI_TaxID=12024;
                                                                                                                                              Viruses; ssRNA positive-strand viruses, unclassified Leviviridae.
                                                                                                                                                               Coat protein.
Bacteriophage PRR1
                                                                                                                                                                                     21-JUL-1986
05-JUL-2004
                                                                                                                                                                                                        21-JUL-1986
                                                                                                    MEDLINE=79148387; PubMed=107028;
                                     3ur. J. Biochem. 94:375-386(1979)
1- FUNCTION: Forms the phage she
IR; A04225; VCBEP1.
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GO:0019028; C:viral capsid; IEA.
GO:0005198; F:structural molecule
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998) to the EMBL/GenBank/DDBJ
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 protein sequencing; RNA-binding.
14535 MW; E7E639E1E50FC612 CRC64;
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Q8VDC2;
01-MAR-2002
01-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
STRALN=BALB/C; TISSUE=Liver;
MEDLINE=22350095; PubMed=12461651;
Kiss H., Darai E., Kiss C., Kost-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=Tmem7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Transmembrane protein
                               01-JUL-1993 (Rel. 26, Last
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Seeligeriolysin precursor
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Listeria seeligeri.
Bacteria; Firmicutes;
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GO:0016021; C:integral to
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AJ428064; CAD20986.1; -.
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Pred. No. 1.2;
17; Mismatches
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Pred. No. 0.
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BIR; S22340; S22340.

HSSP; P19995; 1PF0.

InterPro; IPR001869; Thiol cytolysin.

Pfam; PF01289; Thiol cytolysin; 1.

PRINTS; PR01400; TACYTOLYSIN.

ProDom; PD007062; Thiol cytolysin; 1.

PROSITE; PS00481; THIOL CYTOLYSINS; 1.

CYTOLYSIS; Hemolysis; Lipid-binding; Signal; Toxin.
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Biochim. Biophys. Acta 1130:81-84(1992).
-i- FUNCTION: Sulfhydryl-activated toxin. Is able to lyse cholesterol containing membranes. Can be reversibly inactivated by oxidation. Cholesterol is the receptor for the binding of these toxins to eukaryotic cell membranes.
-i- SIMILARITY: Belongs to the thiol-activated cytolysin family.
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Haas A., Dumbsky M., Kreft J.;
"Listeriolysin genes: complete sequence of ilo from Listeria ivanovii
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530 Seeligeriolysin.
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and repetitive antigen array sectious diseases. The invention non-natural molecular scaffold core particle of a non-natural in and an organiser comprising the organiser is connected to bond. Also disclosed is an east one second attachment		of vaccines for infectious		Lechner F, Sebbel P;			-	-			immunomodulatory; mic; antigen array;						Abg80628 Bacteriop Abr56455 Bacteriop Abr56452 Bacteriop Abr56452 Bacteriop Abu09689 Bacteriop Abu09693 Bacteriop Abr44555 Bacteriop Abr44556 Bacteriop Add24130 Bacteriop Add24131 Bacteriop Add24131 Bacteriop Add82059 Protein f Adj82056 Protein f Add17147 Virus-lik Add17147 Virus-lik Add17147 Virus-lik Add367161 Bacteriop Adj67159 Phage Obe Adj67159 Phage Obe Adj67159 Bacteriop Add52196 Bacteriop Add52196 Bacteriop Add52197 Bacteriop Add52197 Bacteriop Add52198 Bacteriop Add52198 Bacteriop Add52198 Bacteriop Add52198 Bacteriop Add52198 Bacteriop Add52198 Bacteriop

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2001US-0288549P.
2001US-0326998P.
2001US-0331045P.
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Pred. No. 9.6e-70;
Nismatches 0;
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) LECHNER F.
) ORTMANN R.
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) STAUFENBIEL M.
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   LIDAIDQLNPAY 132
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                                                             NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                                 AKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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Pred. No. 9.6e-70;
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RESULT 3 ABR56439

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LIDAIDQLNPAY 132

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ABR56439

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Query Match
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Local Simples 132;
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  Conservative
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100.0%; Pred. No. 9.6e-70;
Live 0; Mismatches 0;
                          Length 132;
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Query Match

100.0%;

Score 663;

B 6 Sequence 132

AA;

cc response against an antigen or a virus-like particle in an animal. (C)
Cc comprises a virus-like particle (VLP) bound to at least one antigen, or a
CC VLP capable of being recognised by the immune system of the animal. Also
CC described: (1) enhancing an immune response against an antigen or a VLP
CC in an animal comprising introducing (C) into the animal; (2) vaccines
CC comprising (C) together with a pharmaceutical diluent, carrier or
CC excipient; (3) immunising or treating an animal comprising administering
CC the vaccine to the animal, or priming or boosting a T cell response in
CC the animal by administering the vaccine; and (4) enhancing anti-viral
CC protection in an animal comprising introducing (C) into the animal. (C)
CC antiallergic, virucide, antibacterial, antiparastic, fungicide,
CC antithyroid, antidiabetic, neuroprotective, nootropic, osteopathic,
CC used for enhancing an immune response against an antigen or a VLP in an
CC used for enhancing anti-viral protection in an animal, or immunising or
CC treating tumours and infectious disease such as viral, bacterial,
CC parastic or fungal infectious diseases such as viral, bacterial,
CC parastic or fungal infectious diseases such as viral, bacterial,
CC parastic or fungal infectious disease, daidetion, graft-versus-bost
CC disease, Crohn's disease, Grave's disease, diabetes, multiple sclerosis,
CC altinumune disease, ACC69833 to ACC69852 and ARR56401 to ARR56500
CC represent sequences used in the exemplification of the present invention.
CC (Updated on 23-OCT-2003 to standardise OS field) cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory; antiithyroid; antiidabetic; neuroprotective; nootropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis; Alzheimer's disease; osteopporosis; rheumatoid arthritis; Disclosure; WPI; 2003-363095/34. Bachmann MF, Storni T, 14-SEP-2001; 2001US-0318967P Bacteriophage Qbeta. Antigen presenting cell; APC; immune response; virus like particle; VLP; Bacteriophage Q-beta coat protein SEQ ID NO:10 23-OCT-2003 28-JUL-2003 The present invention describes a composition (C) for enhancing an immune 16-SEP-2002; 2002WO-IB004252 inflammatory autoimmune disease. ABR56439 composition, useful for enhancing an immune response against an antigen r a virus-like particle, enhancing anti-viral protection in an animal, r immunizing or treating tumors or infectious diseases, e.g. viral CYTOS standard; protein; 132 Page 167-168; (first entry) BIOTECHNOLOGY AG (revised) Lechner 243pp; English য

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                            comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kichney failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of bacteriophage Obeta coat protein CP used in the preparation of the vaccine conjugates of the invention
                                                                                                                                                                                                                                                                                                                        New angiotensin peptide moiety carrier conjugate comprising a carrier as an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WG-1B000166.
19-JUL-2002; 2002US-0396637P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction; concestive heart failure; kidney failure; retinal haemorrhage.
Sequence 132
                                                                                                                                                                                                                                 The invention describes an angiotensin peptide moiety carrier conjugate
                                                                                                                                                                                                                                                                     Claim 15; Page 92; 97pp; English.
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immunostimulatory substance. The immunostimulatory substance is bound to the VLP. Also described: (1) enhancing an immune response in an animal by introducing (C) into the animal; (2) producing (C) for enhancing an immune response in an animal; (3) vaccines comprising (C) together with a
                                                                                                                                                                                                                                                        response
viral inf
                                                                                                                                                                                                                                                                        New compositions comprising immunostimulatory substances packaged into virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, tumors
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(TISS/)
(SCHW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunostimulatory; virus-like particle; bacteriophage; HBV; LCMV; hepatitis B virus; lymphocytic choriomeningitis virus; vaccine; immunostimulant; cytostatic; antiallergic; virucide; antibacterial; immune response; immunisation; allergy; tumour; breast cancer; neuroblastoma; leukaemia; viral disease; influenza; hepatitis; measles; chicken pox; bacterial infection; tuberculosis; pneumonia; syphilis.
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                                                                                                                          The present invention describes a composition (C) for enhancing an immune response in an animal. (C) comprises a virus-like particle (VLP), and an
                                                                                                                                                                                                                                                                                                                                                                                                                                  Maurer P,
Pumpens P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PUMP/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteriophage Q-beta coat protein SEQ ID NO:10.
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MEIJERINK E.
LIPOWSKY G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PUMPENS P.
CIELENS I.
RENHOFA R.
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TISSOT A.
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Cielens :
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                                                                                                                                                                                                      Page 247;
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2002US-0374145P.
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(first entry)
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, Renhofa R,
                                                                                                                                                                                                    English.
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, Bachmann MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                  Lipowsky
7, Storni
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CC (Updated on 23-OCT-2003 to standardise OS field)
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                                                                                      Bachmann M,
                                                                                                                                                                                                     18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-1B000166.
08-JUL-2002; 2002US-039725P.
18-JUL-2002; 2002US-0396590P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteriophage Qbeta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vaccine composition; virus-like particle; core particle; first attachment site; antigen; antigenic determinant; prion PrP; PrP peptide; vaccine; neuroprotective; antiinflammatory; prion disease; Bovine Spongiform Encephalopathy; BSE; Crantyfeldia-Tabak Prioceco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteriophage Qbeta coat protein
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Pred. No. 9.6e-70;
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                                                                                         Renner
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Best Local S
Matches 132
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18-JAN-2002;
21-JAN-2002;
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like or a core particle with at least one first attachment site and at
least one antigen or antigenic determinant that is a prion protein (PrP)
or its dimer, or a PrP peptide, the antigen or antigenic determinant
being bound to the virus-like or core particle. The vaccine of the
invention may have neuroprotective or antiinflammatory activity. The
composition is useful as a medicament or in manufacturing a medicament
for the treatment or prevention of prion diseases. The prion diseases may
include Bovine Spongiform Encephalopathy (BSE) or Creutzfeldt-Jakob
Disease. The present sequence is the amino acid sequence of a coat
protein from a bacteriophage which may be used during the creation of the
vaccine composition of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A vaccine composition for preventing or treating prion diseases (e.g. Creutzfeldt-Jakob Disease) comprises a virus-like particle (e.g. RNA-phage) and at least one prion protein or peptide bound to the virus-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           osteopathic; vaccine; core particle; antigenic determinant; RANKL; bone disease; encephalopathy; immune system stimulation.
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                                                                                                                                            WPI; 2003-441430/41.
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                                                                                                                                                                                                                                                                                                                                                                                                                      ; 2001US-0331045P.
; 2002US-00050902.
; 2002WO-IB000166.
; 2002US-0396635P.
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100.0%; Pred. No. 9.6e-70;
tive 0; Mismatches 0;
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New compositions comprising a core particle and at least one antigen or antigenic determinant, useful for as a vaccine for therapy or prophylaxis

Composition comprising an ordered and repetitive antigen or antideterminant array, useful as a medicament, or for manufacturing

antigenic

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a composition comprising a core particle having at least one first attachment site, and at least one antigen or antigenic determinant having at least one second attachment site. The antigen or antigen or antigenic determinant is a RANKL protein, RANKL tragment or RANKL peptide. The second attachment site is (non-) naturally occurring with the antigen or antigenic determinant, and is capable of association to the first attachment site. The antigen or antigenic determinant and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful as a medicament, or for the manufacture of a medicament for treating bone diseases. The composition is especially useful for as a vaccine for therapy or prophylaxis of bone diseases, particularly mammalian encephalopathies, and for stimulating mammalian immune system. This sequence represents a protein of the invention.
                                                                                                                         07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-1B000166.
19-JUL-2002; 2002US-0396636P.
                                                                                                                                                                                                                                                                                                      antiallergic; antiasthmatic; cytostatic; vaccine; virus-like interleukin; IL-5; IL-13; eotaxin; repetitive antigen array; allergic eosinophilic disease; asthma; Hodgkin's lymphoma.
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                                         WPI; 2003-441518/41.
                                                                                                                                                                                              07-NOV-2002; 2002WO-EP012455.
                                                                                                                                                                                                                                                   WO2003040164-A2
                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                              Virus-like
                                                                                                                                                                                                                                                                                                                                                                                         06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                               ADK17131 standard; peptide; 132 AA.
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Pred. No. 9.6e-70;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                               peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a composition comprising a virus-like particle and at least one antigen, which is a protein or peptide of interleukin (II)-5, II-13 or eotaxin and is bound to the virus-like particle, or a core particle with at least one first attachment site and at least one artigen with at least one second attachment site, where the antigen is a protein or peptide of II-5, II-13 or eotaxin. The second attachment site is capable if an attachment site naturally or not naturally occurring with the cantigen or antigenic determinant. The second attachment site is capable of association to the first attachment site, and where the antigen or antigenic determinant and the core particle interact through the cassociation to form an ordered and repetitive antigen array. The compositions are useful as medicaments, or for manufacturing a medicament or a vaccine for treating allergic eosinophilic diseases, e.g. asthma, or the dogkin's lymphoma and related diseases. This sequence is used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immune response; virus-like particle; immunostimulatory; allergy; chronic disease; chronic viral disease; bacteriophage Qbeta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 132 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteriophage
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     New compositions comprising immunostimulatory substance
                                                                                                                                                                                                                                                          20-JUN-2002; 2002US-0389898P
                                                                                                                                                                                                                                                                                                               20-JUN-2003; 2003WO-EP006541.
                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004000351-A1
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                                                                                        2004-108361/11.
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                                                                                                                                             Renner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VLP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Obeta coat protein virus-like particle
                                                                                                                                                                                                  BIOTECHNOLOGY
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bound to the
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Pred. No. 9.6e-70;
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VLP, and
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        (VLP), an an tigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                an animal comprising a virus-like particle, an immune transmit to the virus-like particle, and an antigen mixed with twirus-like particle. The composition or the vaccine is useful in the manufacture of a pharmaceutical for the treatment of a disease such as allergies, tumours, chronic diseases and chronic viral diseases. The composition is also useful for enhancing an immune responding an animal. This is the amino acid sequence of a bacteriophage Obeta coat protein a virus like particle (VLP) that can be used in the adjuvace of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or o
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                                                                                                                                                                                                                                                                                                                                                        anorectic; core particle; antigenic determinant; ghrelin; P-pilin;
                                                                                                                                                                                                                                                                                                                                                                                                               06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                       ADJ67146;
The invention relates to a new composition comprising: (i) a core particle with at least one first attachment site; and (ii) at least one antigen or antigenic determinant with at least one second attachment
                                                                           New composition comprising a core particle having a first attachment and an antigen or antigenic determinant which is a ghrelin or ghrelin peptide having a second attachment site, useful for treating obesity.
                                                                                                                                                                                                                                                                                                                                                                                   Bacteriophage Qbeta
                                                     Claim 12;
                                                                                                                                                                                                              19-JUL-2002; 2002US-0396638P.
                                                                                                                                                                                                                                       18-JUL-2003; 2003WO-EP007849.
                                                                                                                                                                                                                                                                  29-JAN-2004
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                                                                                                                                                                                                                                                                                                                    Bacteriophage Qbeta
                                                                                                                                                                                                                                                                                                                                              antigenic
                                                                                                                                                           Bachmann MF,
                                                                                                                                                                                    (CYTO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        YLP, useful for enhancing immune response
chronic viral diseases.
                                                                                                                                 2004-132866/13.
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                                                     SEQ
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                                                    4; 175pp; English
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Pred. No. 9.6e-70;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                      for antigen display array
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or antigenic determinant, where the second attachment site is capable of association to the first attachment site, and where the ghrelin or a ghrelin peptide and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful for treating obesity. The repetitive array may form part of a phage or bacterial display array. This peptide corresponds to a Bacteriophage obeta coat protein which can used as part of the repetitive or antigenic array.
Composition comprising a core particle with at least one attachment site, and an antigenic amyloid beta 1-6 peptide, useful for treating diseases such as Alzheimer's disease.
                                                                                                                                                                                                                                             (CYTO-)
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                                                                                                              WPI; 2004-203731/19
                                                                                                                                                                                        Bachmann
                                                                                                                                                                                                                                                                                                                          19-JUL-2002; 2002US-0396639P.
15-MAY-2003; 2003US-0470432P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bacteriophage Qbeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neuroprotective; nootropic; vaccine; amyloid beta 1-6 peptide;
core particle; antigen array; Alzheimer's disease; RNA bacteriophage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacteriophage Qbeta coat protein CP
                                                                                                                                                                                                                                                                                                                                                                                                       18-JUL-2003; 2003WO-EP007864.
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nes 132; Conser
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The invention describes a novel composition particle with at least one attachment site,

comprising a virus-like core and an antigenic amyloid beta

A composition, useful for enhancing or a virus-like particle, enhancing

an immune response against an antigen anti-viral protection in an animal,

Claim 12; SEQ ID NO 4; 184pp; English

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RESULT 12
ABR56440
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1-6 peptide. The new composition comprises: a core particle with one first attachment site; and at least one antigen or antigenic determinant with at least one second attachment site, where the
                                              WPI; 2003-363095/34.
                                                                                                                                                                                                                                                                                      Bacteriophage Qbeta.
                                                                                                                                                                                                                                                                                                                                                       antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus host disease; Grave's disease; diabetes; multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory; antithyroid; antidiabetic; neuroprotective; nootropic; osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteriophage Q-beta coat protein SEQ ID NO:11.
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28-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR56440 standard; protein;
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or antigenic determinant is a Amyloid beta 1-6 peptide, and where
                                                                                                                                                                                  16-SEP-2002;
                                                                                                                                                                                                                    27-MAR-2003
                                                                                                                                                                                                                                                     WO2003024480-A2
                                                                                                                                                14-SEP-2001; 2001US-0318967P.
                                                                                                                                                                                                                                                                                                                                      Alzheimer's disease; osteoporosis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          121
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                                                                                                                  CYTOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                                                                                                                  BIOTECHNOLOGY AG.
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                                                                                                                                                                                                                                                                                                                         disease
                                                                                Lechner
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Pred. No. 9.6e-70;
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Best Local S
Matches 132
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                                                                                                                                                                                                                                                                                         Immunostimulatory; virus-like particle; bacteriophage; HBV; LCMV; hepatitis B virus; lymphocytic choriomeningitis virus; vaccine; immunostimulant; cytostatic; antiallergic; virucide; antibacterial; immune response; immunisation; allergy; tumour; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 328
                                                                               WO2003024481-A2
                                                                                                                                                Bacteriophage Qbeta.
                                                                                                                                                                                                                        neuroblastoma; leukaemia; viral disease; influenza; hepatitis; measles; chicken pox; bacterial infection; tuberculosis; pneumonia; syphilis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteriophage Q-beta coat protein SEQ ID NO:11.
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25-JUL-2003
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Pred. No. 3.6e-69;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                            the VIP. Also described: (1) enhancing an immune response in an animal by introducing (C) into the animal; (2) producing (C) for enhancing an immune response in an animal; (3) vaccines comprising (C) together with a continuous particular or excipient; and (4) immunisting or treating an animal by: (a) administering the vaccine to the animal; (b) priming a T cell response in the animal by administering the vaccine; or (c) boosting a T cell response in the animal by administering the vaccine; or (c) boosting a T cell response in the animal by administering the vaccine; or immune response in an animal, particularly a mammal or unuan.

Specifically, (C) is useful for enhancing a B cell response, a T cell response, or a cytotoxic T-lymphocyte (CTI) response. (C) or a vaccine comprising (C) can also be used for immunising or treating an animal, e.g. humans, sheep, horses, cattle, pigs, dogs, cats, rats, birds, reptiles or fish. (C) is particularly useful in prophylactic or reptiles or fish. (C) is particularly useful in prophylactic or reptiles or specifically, or bacterial infections (e.g. thereat cancers, neuroblastoma, or leukaemia), viral diseases (e.g. influenza, hepatitis, measles or chicken pox), or bacterial infections (e.g. tuberculosis, pneumonia or syphilis). ACC69790 to ACC69815 and ABR44502 to ABR44612 represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 132; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Maurer P, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New compositions comprising immunostimulatory substances packaged into virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, tumors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-SEP-2002; 2002WO-IB004132.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 247-249; 322pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            viral infections.
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22-APR-2002;
                                                                                                                                                                                                                                                                               Sequence 328 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            response in an animal. (C) comprises a virus-like particle (VLP) immunostimulatory substance. The immunostimulatory substance is the VLP. Also described: (1) enhancing an immune response in an introducing (C) into the animal.
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) LIPOWSKY G.
) PUMPENS P.
) CIELENS I.
) RENHOFA R.
                                                                                                                                                                           μ.
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TISSOT A.
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Cielens I,
                                                                                                                                                                                                             Conservative
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2002US-0374145P.
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Pred. No. 3.6e-69;
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R, Bachmann MF,
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27-MAR-2003

ABG94278 standard;

protein;

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                                                                                                                                                                                                                                                                             CC This invention relates to a novel ordered and repetitive antigen array CC used in the production of vaccines for infectious diseases. The invention CC also discloses a composition comprising a non-natural molecular scaffold CC comprising a core particle selected from a core particle of a non-natural CC origin and a core particle of natural origin and an organiser comprising CC at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond. Also disclosed is an CC antigen or antigenic determinant with at least one second attachment CC site, where the antigen or antigenic determinant is amyloid beta peptide (Abetal-42) or its fragment and where the second attachment site is selected from an attachment site not naturally occurring with the antigen CC artitigenic determinant, where the second attachment site is capable of association through at least one non-peptide bond to the first CC attachment site and where the antigen or antigenic determinant and the capable of association through at least one non-peptide bond to the first CC attachment site and where the antigen or antigenic determinant and the CC apable of association through at least one non-peptide bond to the first CC attachment site and where the antigen or antigenic determinant and the CC apable of forming a capsid which comprises mutant Qbeta coat protein CC apable of forming a capsid which comprises mutant Qbeta coat protein CC antiviral, antidiabetic, or hypoglycaemic activities and may be used in immunisation and as a vaccine. The present sequence represents a protein CC antiviral, antidiabetic, or hypoglycaemic activities and may be used in compounds of the invention. (Updated on 06 CC AUG-2003 to Correct OS feld.)
                                                                                          Best Local Similarity Matches 132; Conser
                                                                                                                     Query Match
Best Local
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05-OCT-2001;
07-NOV-2001;
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Piossek C;
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10-DEC-2002
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ilarity 100.0%;
Conservative
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2001US-0326998P.
2001US-0331045P.
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ntiviral; antidiabetic; hypoglycaemic; antigen array;
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                                                                                 Score 663; DB 5;
Pred. No. 3.6e-69;
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The invention relates to a composition comprising: (a) a non-natural molecular scaffold comprising: (i) a core particle selected from: (l) a core particle of a non-natural origin, and (2) a core particle of natural origin; and (ii) an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond, (b) an antigen or antigenic determinant with at least one second attachment site, where the antigen or antigenic determinant is

Disclosure;

Page 393-394; 418pp;

English.

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RESULT 15
ABG80590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Molecular antigen array; vaccine; antigen; antimicrobial; molecular scaffold; amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgE-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus crythematosus; osteoporosis; inflammatory immune disease; lymphadenopathy; multiple sclerosis; immunoproliferative disease; lymphadenopathy; limmunoblastive lymphadenopathy; rheumatoid arthritis; diabetes; infectious disease; factor Xa; rheumatoid arthritis; diabetes; infectious disease; factor Xa;
                                                                    Maurer
Renner
                                                                                                                                                                                                           (CYTO-)
(NOVS )
(MAUR/)
                                                                                                                                                                                                                                                                                                04-MAY-2001;
05-OCT-2001;
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29-NOV-2002
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Molecular antigen array used in
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STAUFENBIEL
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Bachmann
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; 2001US-0288549P.
; 2001US-0326998P.
; 2001US-0331045P.
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CC through the association to form an ordered and repetitive antigen array. The composition is used in client of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context of the context o
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RESULT 2
US-09-948-722-2
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; Sequence 2, Application US/09948722
; Patent No. 6776993
; GENERAL INFORMATION:
; APPLICANT: Kaufmann, Stefan H. E.
; APPLICANT: Hess, Jurgen
; TITLE OF INVENTION: Tuberculosis Va
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; OTHER INFORMATION:
US-09-485-717-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Kaufmann, Stefan
APPLICANT: Hess, Jrgen
TITLE OF INVENTION: Tuberculosis Vaccine
FILE REFERENCE: 16862PUS
CURRENT APPLICATION NUMBER: US/09/485,717
CURRENT FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: EP 97114614.7
PRIOR FILING DATE: 1997-08-22
PRIOR FILING DATE: 1997-08-22
PRIOR APPLICATION NUMBER: PCT/EP98-05109
PRIOR FILING DATE: 1998-08-12
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
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LENGTH: 626
TYPE: PRT
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Best Local Similarity
Matches 31; Conserv
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US-09-256-598-30

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US-09-268-347-48
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PRIOR APPLICATION NUMBER: US 09/485,717
PRIOR PILING DATE: 2000-02-22
PRIOR PELLING DATE: 2000-02-23
PRIOR APPLICATION NUMBER: ECT/EP98/05109
PRIOR APPLICATION NUMBER: EP 97/114,614.7
PRIOR APPLICATION NUMBER: EP 97/114,614.7
PRIOR FILING DATE: 1997-08-23
NUMBER OF SEQ ID NOS: 2
SOPTWARE: Patentin version 3.1
SEQ ID NO 2
LENGTH: 626
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
FEATURE: PRT
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US-09-328-352-4249
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SEQ ID NO 4249
LENGTH: 409
TYPE: PRT
ORGANISM: Acinetobacter baumannii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4249, Application US/09328352
Patent No. 6562958
                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                     -09-328-352-4249
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gary I. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: recombinant nucleic acid OTHER INFORMATION: molecule comprising a domain of Mycobacterium and a phagolysomal OTHER INFORMATION: escape domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                            Local Similarity
                                   117
                                                                        164
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ASKQILDDLPQL 225
                                                                      RTGSN---AIAYLRETGFTNNAALVPENTAKPYSEITF
                                                                                                           RNRKNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALL 116
                                                                                                                                              SLINFTKDVRGGTRLNVNVPRNALISFAVNPVDGTTPI----IAKPNORLTIRDLLAPG 163
                                                                                                                                                                                    TLGNIGKDGKQTLVLN---PR-----GVNPTNGVASLSQAGAVPALEKRVTV-SVSQPS 56
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                                 ASPLLIDAIDQL 128
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Pred.
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No. 0.027;
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                                                                        -EEVMESVKTIAHMLK 213
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RESULT 5
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SOFTWARE: PatentIn Ver. :
SEQ ID NO 49
                                                                                                                                                                                                                                                                     SOFTWARE: P
SEQ ID NO 47
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                                                                                                                        Matches
                                                                                                                                                          Query Match
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Best Local Similarity
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                                                                                                                                           Best
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CURRENT APPLICATION NUMBER: US/09/206,942
CURRENT FILING DATE: 1998-12-08
                                                                                                                                                                                                                                                                                                       EARLIER APPLICATION NUMBER: 09/167,568
EARLIER FILING DATE: 1998-10-07
NUMBER OF SEQ ID NOS: 95
                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Protective Recombinant Haemophilus Influenzae High TITLE OF INVENTION: Molecular Weight Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: LOOSMORE, Sheena M. APPLICANT: Yang, Yan-Ping APPLICANT: Klein, Michel H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/206,942
CURRENT FILING DATE: 1998-12-08
EARLIER APPLICATION NUMBER: 09/167,568
EARLIER FILING DATE: 1998-10-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Yang, Yan-Ping APPLICANT: Klein, Michel H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Protective Recombinant Haemophilus Influenzae High
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                                                                                                                                                                                                                                    TYPE: PRT
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                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Haemophilus influenzae
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                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                        PatentIn Ver.
              PSRN---RKNYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTE
                                                                                     ETVTLGNIGKDGKQTLVLNPRG------VNPTNGVASLSQAGAVPALEKRVTVSVSQ 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ISA 882
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KTLTVGNV--SGNTVTVTANRGALTTLAGSTINGTNGVTTSSQSGEIGGEVTGKTVSVTA
                                                     KTLTVGNV--SGNTVTVTANRGALTTLAGSTINGTNGVTTSSOSGEIGGEVTGKTVSVTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09206942
                                                                                                                          Conservative
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                                                                                                                      11.2%; Score 74; DB 22.8%; Pred. No. 6.3 tive 26; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26; Mismatches
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Pred. No.
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                                                                                                                                                          Length 1079;
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                                                                                                                                                 sequence 4, Application US/08232144
Patent No. 5571695
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
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                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                              APPLICANT: SELBIE, Lisa
APPLICANT: HERZOG, Herbert
APPLICANT: SHINE, John
TITLE OP INVENTION: Human Neuropeptide Y-Y1 Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 383 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cates, Carolyn A.
TITLE OF INVENTION: RHESUS NEUROPEPTIDE Y1 RECEPTOR
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rothwell, Figg, Ernst & Kurz
STREET: 555 13th St, N.W., Suite 701-East
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Gaylo, Paul J.
REGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: P-:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 276-0756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/045,186 FILING DATE: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE:
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5. 6087154
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25.9%;
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Pred. No. 1.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                        DB 3; Length 383;
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Patent No. 5550...
Patent No. 5550...
Parent No. 5550...
Parent No. 5550...
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                                                                                                                                COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC LDOS/MS-DOS

COPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: ERNST, Barbara G
REGISTRATION NUMBER: 30,377
REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
   CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: 811
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OPERATING SYSTEM:
                                                                       FILING DATE:
                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                          ADDRESSEE:
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                                                                                        APPLICATION NUMBER:
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SYSTEM: PC-DOS/MS-DOS
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25.9%; Pred. No. 1.4;
tive 12; Mismatches
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of the Americas
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SOFTWARE: PatentIn Release #1.0, Version #1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      138 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16 KQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
                                                                                                                                                                                                                                                                                                                                                                              k, Richard L.

DNA ENCODING A HUMAN NEUROPEPTIDE
Y/PEPTIDE YY/PANCREATIC POLYPEPTIDE
RECEPTOR (Y4) AND USES THEREOF
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US-10-013-846-4
; Sequence 4, Application US/10013846
; Patent No. 6566367
; GENERAL INFORMATION:
; APPLICANT: Bakthavatchalam, Rajagor
; APPLICANT: Blum, Charles A
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US-09-200-673-15
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                                                                                                                     RESULT 10
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US-09-200-673-15
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Matches
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SEQ ID NO 15
LENGTH: 384
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Patent No. 6316203
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gerald, Christophe P.G.
APPLICANT: Weinshank, Richard L.
APPLICANT: Walker, Mary W.
APPLICANT: Branchek, Theresa
TITLE OF INVENTION: Methods of Modifying Feeding Behavior, Compounds Useful
TITLE OF INVENTION: in Such Methods, and DNA Encoding A Hypothalamic
TITLE OF INVENTION: Atypical Neuropeptide Y/Peptide YY Receptor (Y5)
FILE REFERENCE: 46166-BZ/JPW
                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 38
TYPE: PRT
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
TELEPAX: (212) 391-0526
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                                                                                                                                                                                                                                                                                           16 KQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
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                     Rajagopal
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Pred. No. 1.4;
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Pred. No. 1.4;
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RESULT 11
US-09-708-392-9
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                                                                                                             PRIOR FILING DATE: 2000-07-11
PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2000-07-27
PRIOR PRIOR PRIOR DATE: 2000-07-27
NUMBER OF SEQ ID NOS: 20
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Pa
SEQ ID NO 9
LENGTH: 384
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APPLICANT: Elliott, Richard L
APPLICANT: Hammond, Marlys
TITLE OF INVENTION: Spiro[isobenzofuran-1,4'-piperidin]-3-ones and
TITLE OF INVENTION: 3H-spiroisobenzofuran-1,4'-piperidines
FILE REFERENCE: NOO.2001
CURRENT APPLICATION NUMBER: US/10/013,846
CURRENT FILING DATE: 2001-12-11
PRIOR APPLICATION NUMBER: US 60/254,990
PRIOR FILING DATE: 2000-12-12
NUMBER OF SEQ ID NOS: 17
SOPTWARE: Patentin version 3.1
SEQ ID NO 4
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Patent No. 6734186
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PRIOR APPLICATION NUMBER: US (
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CURRENT FILING DATE: 2000-11-08
PRIOR APPLICATION NUMBER: GB 9926437.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-0
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 2000-0
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PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/175,161
APPLICATION NUMBER: US 60/175,161
APPLICATION NUMBER: US 60/175,161
FILING DATE: 2000-01-07
APPLICATION NUMBER: US 60/192,962
APPLICATION NUMBER: US 60/192,962
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                                                                                    PatentIn Ver.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          De Lombaert, Ste
Hutchison, Alan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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; TYPE: PRT; ORGANISM: Homo sapiens US-09-708-392-9
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RESULT 13
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Best Local Similarity
Matches 28; Conserv
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GENERAL INFORMATION
APPLICANT: Claes
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Best Local Similarity 25.9%;
Matches 28; Conservative 1:
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MEDIUM TYPE: floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: MS-DOS
SOFTWARE: Microsoft Word 4.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/0:
FILING DATE: 1930527
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
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TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: GEOTGE M. Yahwak
REGISTRATION NUMBER: 26,824
REFERENCE/DOCKET NUMBER: CR
                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                       TYPE: AMINO ACID
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                                                                                                                                                           16 KQTLVLNPRGVNPTN-----GVA----SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
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                                                                                    IQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAAL 115
                                                                                                                       RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN--
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(203)268-1951
(203)268-1951
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linear
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25.9%; Pred. No. 1.4;
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Receptor of the Y1-Type and
Anitsense Oligonucleotides
Thereto Which Inhibit Vasoconstriction
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                                                    VTLDAYKDKYVCFDQFPSDSHRLSYTTLLLVL 218
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Pred. No. 1.4;
                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                             33; Indels 35;
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                                                                                                                                                                                                                              Length 384;
                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                Sequence 3, Application PC/TUS9514377 GENERAL INFORMATION:
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                       APPLICANT: STRADER, CATHERINE D.
APPLICANT: CASCIERI, MARGARET A.
APPLICANT: MACHELL, DOUGLAS J.
TITLE OF INVENTION: MODIFIED NEUROPEPTIDE Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 25,147
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION IMFORMATION:
TELEPHONE: (732) 594-1348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: CASCIERI, MARGARET APPLICANT: MACNEIL, DOUGLAS J. TITLE OF INVENTION: MODIFIED N
                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: MARY A. APPOLLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US 0
FILING DATE: 07-NOV-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                          NUMBER OF SEQUENCES: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                       ADDRESSEB: MARY A. AFFURET: 126 EAST LINCOLN AVENUE
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CLASSIFICATION:
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                                                     CITY: RAHWAY
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                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: NICHOLSON,
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                                                                                                                                                                                                                                                                                                                                 190 --
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07065-0900
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                                  NEW JERSEY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11.1%; Score 73.5; D
25.9%; Pred. No. 1.6;
                                                                                                                                                                                                                                                                                                                                 ----VTLDAYKDKYVCFDQFPSDSHRLSYTTLLLVL 221
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US-08-397-411-13; Sequence 13, A
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Best Local S
              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/397,411
FILING DATE: 01-MAR-1995
CLASSIFICATION 124
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/859,583
FRIOR APPLICATION UNMBER: US 07/859,583
FILING DATE: 27-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/335,017
APPLICATION NUMBER: US 08/335,017
FILING DATE: 07-NOV-1994
ATTORNEY/AGENT INFORMATION:
NAME: APPOLLINA, MARY A.
REGISTRATION NUMBER: 34,087
REFERENCE/DOCKET NUMBER: 19339Y POTELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (908) 594-3462
TELEFAX: (908) 594-4720
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/14377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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MEDIUM TYPE: Floppy disk
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                            TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                         STREET: One Market a CITY: San Francisco STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               PLICANT: Tso, J. Yun
TLE OF INVENTION: Bispecific Antibody Effective to Treat
TLE OF INVENTION: B-Cell Lymphoma and Cell Line
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1 Similarity 25.9%; Pred. No. 1.6;
28; Conservative 12; Mismatches
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Search completed: January 4, 2005, 09:35:40
Job time : 11.3622 secs
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                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Smith, William M. REGISTRATION NUMBER: 30,223 REFERENCE/DOCKST NUMBER: 01: TELECOMMUNICATION INFORMATION: TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
TOPOLOGY: lir
MOLECULE TYPE:
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                                                                       238 GLTDTLQAETD --
                                                                                                                                          179 SSGLYSLSSVVTVPSSSSLGTQTYICNVNHKPSNTKVDKKVE-PKSCDKTHTCPPCKCPAG 237
                                                                                                       83 --SVTROAYADVTFSFTQYSTDEERAFVRTELAALL 116
                                                                                                                                                                            29 TNGVASLSQAGAVPALEKRVTVSVSQPSRNRKNYKYQVKIQNPTACTANGSCDP----- 82
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amino acid
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Minimum
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Maximum Match 100%
Listing first 45 s
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Perfect score:
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seq length:
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1: /cgn2 6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2 6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

3: /cgn2 6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

4: /cgn2 6/ptodata/1/pubpaa/US06 PUBCOMB.pep:*

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6: /cgn2 6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

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11: /cgn2 6/ptodata/1/pubpaa/US09 PUBCOMB.pep:*

12: /cgn2 6/ptodata/1/pubpaa/US09 NEW PUB.pep:*

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18: /cgn2 6/ptodata/1/pubpaa/US10 NEW PUB.pep:*

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                                                                                                                                                                                                                                                                                              Query
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Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                           Length
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   US-10-243-739-10
US-10-244-65-10
US-10-289-454-10
US-10-050-902-159
US-10-050-908-159
US-10-346-190-10
US-10-346-190-10
US-10-369-456-10
US-10-622-064-3
US-10-622-124-4
US-10-623-739-11
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1450.557 Million cell updates/sec
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Sequence 10, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 159, App
Sequence 159, App
Sequence 10, Appl
Sequence 1, Appl
Sequence 1, Appl
Sequence 3, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 11, Appl
Sequence 11, Appl
                                                                                                                                                                                                                                                                     APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Lechner, Franziska
FITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of
FITLE OF INVENTION: Inwivo Activation of Antigen Presenting Cells for Enhancement of
FITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
FILE REFERENCE: 1700.0210001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 10
LENGTH: 132
TYPE: PRT
ORGANISM: Bacteriophage Q-beta
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Matches 132
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Similarity 100.0%; Pred. No. 1.6e-66.
32; Conservative 0; Mismatches 0.
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                                                                                                                                           AKLETYTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRYTVSVSQPSRNRK 60
                                             NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL
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US-10-050-989-817-1
US-10-346-190-11
US-10-465-811-2
US-10-622-064-4
US-10-622-087-5
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22, Appl
23, Appl
24, Appl
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Result No.

60

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APPLICANT: Cielens, Indulis
APPLICANT: Cielens, Indulis
TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Particl
TITLE OF INVENTION: Method of Preparation and Use
FILE REFERNCE: 1700.0220001
CURRENT APPLICATION NUMBER: US/10/244,065
CURRENT APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR FILING DATE: 2002-04-22
PRIOR APPLICATION ON THE COMMENT OF SEQ ID NO: 03-01-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 10
LENGTH: 132
TYPE: PRI
ORGANITOTI
                                                                                                                                                                                                                                                                                                                                                             US-10-289-454-10; Sequence 10, Application US/10289454; Publication No. US20030157479A1
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Best Local Similarity
APPLICANT: Jennings, Gary
APPLICANT: Jennings, Gary
APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
CURRENT APPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-01-19
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
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; TYPE: PRT ; ORGANISM: Bacteriophage Q-beta US-10-289-454-10
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APPLICANT: Maurer, Patrick
APPLICANT: Maurer, Patrick
APPLICANT: Lechner, Franziska
APPLICANT: Sebbel, Peter
APPLICANT: Piossek, Christine
TITLE OF INVENTION: Molecular Antigen Array
FILE REFERENCE: 1700.0190004
CURRENT APPLICATION NUMBER: US/10/050,902
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR APPLICATION NUMBER: US 60/268,549
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-05-04
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Best Local Similarity 100.0%;
Matches 132; Conservative 0
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 159
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APPLICANT: Renner, Wolfgang A.
APPLICANT: Bachmann, Martin
APPLICANT: Tissot, Alain
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 10
LENGTH: 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
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TYPE: PRT
ORGANISM: Bacteriophage Q Beta
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                                                                                      61 NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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Pred. No. 1.6e-66;
); Mismatches 0;
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LIDAIDQLNPAY 132

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CURRENT APPLICATION NUMBER: US/10/050,898
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-05-04
PRIOR PILING DATE: 2001-10-05
PRIOR PILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR PILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 350
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 159
APPLICANT: Bachmann, Martin
APPLICANT: Bachmann, Martin
APPLICANT: Renner, Patrick
APPLICANT: Pellicioli, Erica
APPLICANT: Renner, Wolfgang A.
TITLE OF INVENTION: Prion Protein Carrier-Conjugates
FILE REFERENCE: 1700.0290003
CURRENT APPLICATION NUMBER: US/10/346,190
CURRENT FILING DATE: 2003-01-17
PRIOR APPLICATION NUMBER: 60/396,590
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,725
PRIOR APPLICATION NUMBER: 60/393,898
PRIOR APPLICATION NUMBER: 60/389,898
PRIOR FILING DATE: 2002-06-20
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; Sequence 10, Application US/10346190
; Publication No. US20030219459A1
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; ORGANISM: Bacteriophage Q Beta
US-10-050-898-159
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US-10-050-898-159
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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Luond, Rainer
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Sebbel, Peter
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Tissot, Alain
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Pred. No. 1.6e-66;
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RESULT 8
US-10-289-456-10
; Sequence 10, Application US/10289456
; Publication No. US20040033211A1
; GENERAL INFORMATION:
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; TYPE: PRT
; ORGANISM: Bacteriophage Q-beta
US-10-465-811-1
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PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR FILING DATE: 2002-01-18
NUMBER OF SEQ ID NOS: 164
SOFTWARE: PatentIn version 3.1
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/10465811 Publication No. US20040005338A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.2 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 132; Conservative 0
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Best Local (
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APPLICANT: RENNER, WOLFGANG A
TITLE OF INVENTION: PACKAGED VIRUS-LIKE PARTICLES
TITLE OF INVENTION: METHOD OF PREPARATION AND USE
FILE REFERENCE: 1700.0290004
CURRENT APPLICATION NUMBER: US/10/465,811
CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: US 60/389,898
PRIOR PILING DATE: 2002-06-20
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                                                                                                                                                                                                                                                   NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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Pred. No. 1.6e-66
); Mismatches 0
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Bachmann, Martin

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APPLICANT: Bachmann, Martin F
APPLICANT: Bachmann, Martin F
TITLE OF INVENTION: Hapten-Carrier Conjugates and Uses Thereof
FILE REFERENCE: 1700.0300001
CURRENT APPLICATION NUMBER: US/10/622,064
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,575
PRIOR FILING DATE: 2002-07-18
NUMBER OF SEQ ID NOS: 33
SOFTWARE: Patentin version 3.2
SEQ ID NO 3
LENGTH: 132
TYPEE: PRT
CORGANISM: Bacteriophage Obeta
US-10-622-064-3
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PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
PRIOR FILING DATE: 2001-11-07
PRIOR FILING DATE: 2001-11-07
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SEQ ID NO 10
                                                                                                                                                                                         Query Match
Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/289,456
CURRENT FILING DATE: 2002-11-07
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TYPE: PRT
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APPLICANT: Spohn, Gunther
TITLE OF INVENTION: Antigen Arrays for Treatment of Bone Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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                    61 NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                    AKLETYTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                                                                                                                                                100.0%; Score 663; DB 15; ilarity 100.0%; Pred. No. 1.6e-66; Conservative 0; Mismatches 0;
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APPLICANT: Bachmann, Martin F
APPLICANT: Tissot, Alain
APPLICANT: Tissot, Rainer
APPLICANT: Ortmann, Rainer
APPLICANT: Lound, Rainer
APPLICANT: Staufenbiel, Matthias
APPLICANT: Frey, Peter
APPLICANT: Frey, Peter
TITLE OF INVENTION: Amyloid Beta 1-6 Antigen Arrays
FILE REFERENCE: 1700.0350002
CURRENT APPLICATION NUMBER: US/10/622,087
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,639
PRIOR FILING DATE: 2002-07-19
PRIOR FILING DATE: 2003-05-15
PRIOR FILING DATE: 2003-05-15
NUMBER OF SEQ ID NOS: 93
NUMBER OF SEQ ID NOS: 93
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APPLICANT: Fulurija, Alma
APPLICANT: Fulurija, Alma
TITLE OF INVENTION: Ghrelin-Carrier Conjugates
FILE REFERENCE: 1700.0340001
CURRENT APPLICATION NUMBER: US/10/622,124
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US 60/396,638
PRIOR FILING DATE: 2002-07-19
NUMBER OF SEQ ID NOS: 146
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US-10-622-124-4
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                                                                                   ; TYPE: PRT ; ORGANISM: Bacteriophage Q-beta US-10-622-087-4
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Publication No. US20040141984A1
GENERAL INFORMATION:
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Best Local Similarity 100.0%;
Matches 132; Conservative 0
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Publication No. US20040076645A1
Query Match
Best Local Similarity
Matches 132; Conserv
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      Conservative
  100.0%; Score 663; DB 16;
100.0%; Pred. No. 1.6e-66;
... Mismatches 0;
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Pred. No. 1.6e-66;
); Mismatches 0;
                                            Length 132;
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      Gaps
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RESULT 13
US-10-244-065-11
; Sequence 11, Application US/10244065
; Publication No. US2003009966BA1
; GENERAL INFORMATION:
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US-10-243-739-11
J Sequence 11, Application US/10243739
Publication No. US20030091593A1
GENERAL INFORMATION:
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; ORGANISM: Bacteriophage Q-beta
US-10-243-739-11
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                                                                                                                                                   APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Maurer, Patrick
APPLICANT: Tissot, Alain
APPLICANT: Schwarz, Katrin
APPLICANT: Schwarz, Edwin
APPLICANT: Lipowsky, Gerad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bachmann, Martin F.
APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Lechner, Franziska
TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of
TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
TILE REFERENCE: 1700.0210001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
LENGTH: 328
TVDE: DDT
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Best Local Similarity
APPLICANT: Schwarz, Katrin
APPLICANT: Meijerink, Edwin
APPLICANT: Lipowsky, Gerad
APPLICANT: Lipowsky, Gerad
APPLICANT: Pumpens, Paul
APPLICANT: Cielens, Indulis
APPLICANT: Renhofa, Regina
TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like |
FILE REFERENCE: 1700.0220001
CURRENT APPLICATION NUMBER: US/10/244,065
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; LENCTH: 329
; TYPE: PRT
; ORGANISM: Bacteriophage Q-beta
US-10-289-454-11
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; ORGANISM: Bacteriophage Q-beta
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CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR PPLICATION NUMBER: US 10/050,902
PRIOR PILING DATE: 2002-01-18
PRIOR PPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR PILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 386
SOFTWARE: PatentIn version 3.2
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Matches
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APPLICANT: Bachmann, Martin
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LENGTH: 328
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PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOPTWARE: Patentin version 3.1
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PRIOR APPLICATION NUMBER: 60/374,145
PRIOR FILING DATE: 2002-04-22
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TITLE OF INVENTION: Antigen Arrays for Treatments
FILE REFERENCE: 1700.0360001
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62 NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL AKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60 NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL AKLETVTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 61

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US-10-050-902-217

Sequence 217, Application US/10050902

Publication No. US20030175290A1

GENERAL INFORMATION:

APPLICANT: Renner, Wolfgang A.

APPLICANT: Bachmann, Martin

APPLICANT: Hachmann, Martin

APPLICANT: Sebbel, Peter

APPLICANT: Pioseek, Christine

TITLE OF INVENTION: Molecular Antigen Array

FILE REFERENCE: 1700.019004

CURRENT APPLICATION NUMBER: US(10/050,902)

CURRENT FILING DATE: 2001-01-19

PRIOR APPLICATION NUMBER: US 60/262,379

PRIOR APPLICATION NUMBER: US 60/262,379

PRIOR APPLICATION NUMBER: US 60/288,549

PRIOR APPLICATION NUMBER: US 60/326,998

PRIOR APPLICATION NUMBER: US 60/326,998

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PRIOR APPLICATION NUMBER: US 60/331,045

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   19-JAN-2001;
04-MAY-2001;
05-OCT-2001;
07-NOV-2001;
This invention relates to a novel ordered and repetitive antigen array used in the production of vaccines for infectious diseases. The invention also discloses a composition comprising a non-natural molecular scaffold comprising a core particle selected from a core particle of a non-natural origin and a core particle of natural origin and an organiser comprising at least one first attachment site, where the organiser is connected to the core particle by at least one covalent bond, Also disclosed is an antigen or antigenic determinant with at least one second attachment
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Piossek C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unidentified bacteriophage.
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; 2001US-0288549P.
; 2001US-0326998P.
; 2001US-0331045P.
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Result No.

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Query Match

Length

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Description

pQB259

ABG94319 ABG80631

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Minimum

Total number Searched:

of hits satisfying chosen parameters:

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Scoring table:

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GenCore version (c) 1993 - 2005

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                          19-JAN-2001;
04-MAY-2001;
05-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the antigen or antigenic determinant, where the second attachment site is capable of association through at least one non-peptide bond to the first attachment site and where the antigen or antigenic determinant and the scaffold interact through the association to form an ordered and repetitive antigen array. The invention also comprises a coat protein capable of forming a capsid which comprises mutant Obeta coat proteins having an amino acid sequence selected from five amino acid sequences fully defined in the specification. The compounds of the invention may have antimicrobial, antiallergic, immunomodulatory, cytostatic, antiviral, antidiabetic, or hypoglycaemic activities and may be used in immunisation and as a vaccine. The present sequence represents a protein sequence used to create the compositions of the invention. (Updated on 29 -AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecular antigen array; vaccine; antigen; antimicrobial; molecular scaffold; amyloid beta, Abeta 1-42; influenza; graft versus host disease; IgE-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus erythematosus; osteoporosis; influence disease; systemic lupus erythematosus; osteoporosis;
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                                                             07-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteriophage
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The invention relates to a composition comprising: (a) a non-natural completular scaffold comprising: (i) a core particle of a non-natural coigin; and (2) a core particle of a non-natural origin; and (2) a core particle of natural coigin; and (2) a core particle of natural coigin; and (2) a core particle of natural corigin; and (2) a core particle of natural coigin; and (3) a core particle by at least conjugation or site, where the core particle by at least cone second attachment site, where the antigen or antigenic determinant with at least cone second attachment site, where the antigen or antigenic determinant is amyloid beta peptide (Abeta 1-42) or its fragment, and where the second cattachment site is selected from. (i) an attachment, and where the second determinant, where the second attachment site is capable of association composition at least one non-peptide bond to the first attachment site; and where the antigen or antigenic determinant and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is used in condered and repetitive antigen array. The composition is anaphylaxis, adult respiratory distress syndrome (ARDS), Crohn's disease, allergic asthma, cause in ymphodenopathy, immunoblastive lymphadenopathy, cause in the particle of a condered and infectious diseases. The antigens are modified to posses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 18; Page 403; 418pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecular antigen array used in the
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MAURER P.
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ORTMANN R.
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STAUFENBIEL M.
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Bachmann
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Sebbel
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Best Local
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                             NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                 ARLETYTLGNIGKDGROTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                                                                                                                100.0%; Score 664; DB 5 ilarity 100.0%; Pred. No. 2e-71; Conservative 0; Mismatches
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RESULT 3
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121

LIDAIDQLNPAY 132

ABR56456 standard; protein; 132

antithyroid; antidiabetic; neuroprotective; nootropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis; Alzheimer's disease; osteoporosis; rheumatoid arthritis;

cytostatic; virucide; antibacterial; antiparasitic; fungicide; antiallergic; immunosuppressive; antiaddictive; antiinflammatory;

Antigen presenting cell; APC; immune response; virus like particle;

Bacteriophage Q-beta coat protein mutant SEQ ID

NO:27.

VLP;

28-JUL-2003 ABR56456;

(first entry)

Bacteriophage Qbeta.

inflammatory autoimmune disease.

WO2003024480-A2

14-SEP-2001; 2001US-0318967P. 16-SEP-2002; 2002WO-IB004252.

(CYTO-) CYTOS BIOTECHNOLOGY AG

Storni 'n

Lechner F;

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response against an antigen or a virus-like particle in an animal. (C) comprises a virus-like particle (VLP) bound to at least one antigen, or a VLP capable of being recognised by the immune system of the animal. Also described: (1) enhancing an immune response against an antigen or a VLP in an animal comprising introducing (C) into the animal; (2) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; (3) immunising or treating an animal comprising administering the vaccine to the animal, or prining or boosting a T cell response in the animal by administering the vaccine; and (4) enhancing anti-viral protection in an animal comprising introducing (C) into the animal. (C) antiallergic, virucide, antibacterial, antiparasitic, fungicide, antithyroid, antidiabetic, neuroprotective, nootropic, osteopathic, antithrematic and antiarthritic activities. (C) or the vaccines can be antifered.
                              used for enhancing an immune response against an antigen or a VLP in an animal, enhancing anti-viral protection in an animal, or immunising or treating tumours and infectious diseases such as viral, bacterial, parasitic or fungal infections. The vaccine compositions are also useful for preventing or treating allergies, drug addiction, graft-versus-host disease, Crohn's disease, Grave's disease, diabetes, multiple sclerosis, Alzheimer's disease, osteoporosis, rheumatoid arthritis, or inflammatory autoimmune disease. ACC69838 to ACC69852 and ABR56401 to ABR56509 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A composition, useful for enhancing an immune response against an antigen or a virus-like particle, enhancing anti-viral protection in an animal, or immunizing or treating tumors or infectious diseases, e.g. viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a composition (C)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             243pp; English.
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                                                                                     The invention describes an angiotensin peptide moiety carrier conjugate comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kidney failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta coat protein used in the preparation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New angiotensin peptide moiety carrier conjugate comprising a carrier and an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-OCT-2001; 2001US-0326998P.
07-NOV-2001; 2001US-03310459.
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-1B000166.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteriophage Obeta; coat protein; hypotensive; cerebroprotective; cardiant; nephrotropic; ophthalmological; immunostimulant; vaccine; angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke; infarction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 20; Page 94; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or infarction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-430264/40
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; 2002US-00050902.
; 2002WO-IB000166.
; 2002US-0396637P.
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Sequence 132 A

Query Match Best Local Similarity 100.0%; Score 664; DB 6 Pred. No. 2e-71; 6; Length 132;

ARLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60

Matches 132;

Local Similarity

100.0%; ilarity 100.0%; Conservative

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Score 664; DB 6 Pred. No. 2e-71; Mismatches

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Length 132; Indels

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Query Match

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RESULT 5
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The present invention describes a composition (C) for enhancing an immune response in an animal. (C) comprises a virus-like particle (VLP), and an immunostimulatory substance. The immunostimulatory substance is bound to the VLP. Also described: (1) enhancing an immune response in an animal by introducing (C) into the animal; (2) producing (C) for enhancing an immune response in an animal; (3) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; and (4) immunising or
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Maurer P,
Pumpens P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunostimulatory; virus-like particle; bacteriophage; HBV; LCMV; hepatitis B virus; lymphocytic choriomeningitis virus; vaccine; immunostimulant; cytostatic; antiallergic; virucide; antibacterial; immune response; immunisation; allergy; tumour; breast cancer; neuroblastoma; leukaemia; viral disease; influenza; hepatitis; measles; chicken pox; bacterial infection; tuberculosis; pneumonia; syphilis.
                                                                                                                                                                                                                                                                                             New compositions comprising immunostimulatory substances packaged into virus-like particles, useful as a vaccine for enhancing an immune response in animals, e.g. for treating or preventing allergies, tumors
                                                                                                                                                                                                                        Disclosure;
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22-APR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SCHWARZ K.
MEIJERINK E.
LIPOWSKY G.
PUMPENS P.
CIELENS I.
RENHOFA R.
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TISSOT A.
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Cielens I,
                                                                                                                                                                                                                     Page 262; 322pp; English.
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                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vaccine composition; virus-like particle; core particle; first attachment site; antigen; antigenic determinant; prion PrP; PrP peptide; vaccine; neuroprotective; antiinflammatory; prion disease; Bovine Spongiform Encephalopathy; BSE;
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                                                                                                                                                                                                                                        18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002WO-1B000166.
08-JUL-2002; 2002US-039725P.
18-JUL-2002; 2002US-0396590P.
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priming a T cell respon
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Pred. No. 2e-71;
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Matches
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18-JAN-2002;
21-JAN-2002;
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like or a core particle with at least one first attachment site and at
least one antigen or antigenic determinant that is a prion protein (PrP)
or its dimer, or a PrP peptide, the antigen or antigenic determinant
being bound to the virus-like or core particle. The vaccine of the
invention may have neuroprotective or antiinflammatory activity. The
composition is useful as a medicament or in manufacturing a medicament
for the treatment or prevention of prion diseases. The prion diseases may
include Bovine Spongiform Encephalopathy (BSS) or Creutzfeldt-Jakob
Disease. The present sequence is the amino acid sequence of a mutant coat
protein from bacteriophage Obeta which may be used during the creation of
the vaccine composition of the invention.
New compositions comprising a core particle and at least one antigen or antigenic determinant, useful for as a vaccine for therapy or prophylaxis of bone diseases, particularly mammalian encephalopathies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein for RANKL antigen array to treat bone disease
                                                                                                                          WPI; 2003-441430/41.
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                                                                                                                                                                                                                                            (CYTO-) CYTOS BIOTECHNOLOGY AG
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2002US-00050902.
2002WO-IB000166.
2002US-0396635P.
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Best Local Similarity
Matches 132; Conserv
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Composition comprising an ordered and repetitive antigen or antigenic determinant array, useful as a medicament, or for manufacturing a medicament for treating allergic eosinophilic diseases, e.g. asthma,
                                                                                                                                                                                                                                                                                                                                                                antiallergic; antiasthmatic; cytostatic; vaccine; interleukin; IL-5; IL-13; eotaxin; repetitive antiallergic eosinophilic disease; asthma; Hodgkin's
                                                                                                                                                         07-NOV-2001; 2001US-0331045P.
18-JAN-2002; 2002US-00050902.
21-JAN-2002; 2002US-1B000166.
19-JUL-2002; 2002US-0396636P.
                                                                                                                                                                                                                                      07-NOV-2002; 2002WO-EP012455.
                                                                                                                                                                                                                                                                                                     WO2003040164-A2
                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                              Virus-like particle
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                                                               2003-441518/41.
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                                                                                                                                                                                                                                                                                                                                                                                                                              repetitive antigen array peptide
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Pred. No. 2e-71;
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n's|lymphoma.
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Disclosure; SEQ ID NO Hodgkin's lymphoma

27; 245pp; English

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Best Local Simi
Matches 132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immune response; virus-like particle; immunostimulatory; allergy; tumour;
chronic disease; chronic viral disease; bacteriophage Qbeta;
coat protein; VLP; adjuvant; mutant; mutein.
   New compositions comprising a virus-like particle (VLP), an immunostimulatory substance bound to the VLP, and an antigen mixed the VLP, useful for enhancing immune response or for treating e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 132 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                 20-JUN-2003; 2003WO-EP006541
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nilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             virucide; immunostimulant; vaccine;
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Pred. No. 2e-71;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADJ67163
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The invention relates to a new composition comprising: (i) a core particle with at least one first attachment site; and (ii) at least one antigen or antigenic determinant with at least one second attachment
                                                                                                                                                           New composition comprising a core particle having a first attachment site and an antigen or antigenic determinant which is a ghrelin or ghrelin peptide having a second attachment site, useful for treating obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bacteriophage Qbeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Phage Qbeta
                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                 WPI; 2004-132866/13.
                                                                                                                                                                                                                                                                                                                                        Bachmann MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-2002; 2002US-0396638P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-JUL-2003; 2003WO-EP007849.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               anorectic; core particle;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDBBRAFVRTBLAALLASPL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  coat protein mutant Qbeta 259 for antigen display array
                                                                                                                SEQ ID NO
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                                                                                                                                                                                                                                                                                                                                     Fulurija
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antigenic determinant; ghrelin; P-pilin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            132 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 8;
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Best Local Similarity
Matches 132; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuroprotective; nootropic; vaccine; amyloid beta 1-6 peptide; core particle; antigen array; Alzheimer's disease; RNA bacteriophage; coat protein; CP; mutant; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for treating obesity. The repetitive array may form part of a phage or bacterial display array. This peptide corresponds to a Bacteriophage Obeta coat protein mutant which can used as part of the repetitive or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or antigenic determinant, where the second attachment site is capable of association to the first attachment site, and where the ghrelin or a ghrelin peptide and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage Obeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteriophage Qbeta coat protein mutant K2R K16R
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                                                      WPI; 2004-203731/19
                                                                                                                                    Bachmann MF,
                                                                                                                                                                                                                                                                                  19-JUL-2002;
15-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                18-JUL-2003; 2003WO-EP007864.
                                                                                                                                                                                                 (NOVS ) NOVARTIS PHARMA
                                                                                                                                                                                                                           (CYTO-) CYTOS BIOTECHNOLOGY AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; protein;
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2003US-0470432P.
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                                                                                                                                          Tissot
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Wild type Lys substituted by Arg"
                                                                                                                                       Þ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Wild type
                                                                                                                                                                                                 ĀG.
                                                                                                                                          Ortmann
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Pred. No. 2e-71;
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                                                                                                                                       Lueoend R,
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Composition comprising a core particle with at least one attachment site,

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RESULT 12
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                                                                                    19-JAN-2001;
04-MAY-2001;
05-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or antigenic determinant is a Amyloid beta 1-6 peptide, and where the second attachment site comprises: an attachment site not naturally occurring with the antigen or antigenic determinant; or an attachment site naturally occurring with the antigen or antigenic determinant. The second attachment site is capable of association to the first attachment site and the beta 1-6 peptide and the core particle interact through the association to form an ordered and repetitive antigen array. The composition is useful for the manufacture of a medicament for treating Alzheimer's disease and related diseases. This is the amino acid sequence of an RNA bacteriophage Obeta coat protein mutant that can be used in the preparation of the compositions and vaccines of the invention.
                                                                                                                                                                                                                                                                         Human; mouse; rat; antimicrobial; antiallergic; immorptostatic; antiviral; antidiabetic; hypoglycaemic; vaccine; infectious disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention describes a novel composition comprising a virus-like core particle with at least one attachment site, and an antigenic amyloid beta 1-6 peptide. The new composition comprises: a core particle with at least one first attachment site, and at least one antigen or antigenic determinant with at least one second attachment site, where the antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and
Renner WA,
Piossek C;
                                                                                                                                                     21-JAN-2002; 2002WO-IB000166.
                                                                                                                                                                                                                  WO200256905-A2
                                                                                                                                                                                                                                             unidentified bacteriophage.
                                                                                                                                                                                                                                                                                                                                       pQB251 protein.
                                                                                                                                                                                                                                                                                                                                                                    29-AUG-2003
10-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                               ABG94320 standard; protein; 132 AA
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                                            (CYTO-) CYTOS BIOTECHNOLOGY AG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    an antigenic amyloid beta 1-6 peptide, useful for treating as Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             121 LIDAIDQLNPAY 132
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                                                                         ; 2001US-0262379P.
; 2001US-0288549P.
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; 2001US-0331045P.
               Bachmann
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Pred. No. 2e-71;
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Best Local Sim
Matches 131;
                                     molecular scaffold; amyloid beta; Abeta 1-42; influenza; graft versus host disease; IgB-mediated allergic reaction; anaphylaxis; adult respiratory distress syndrome; ARDS; Crohn's disease; allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma; Grave's disease; systemic lupus erythematosus; osteoporosis; inflammatory immune disease; mysthematosus; osteoporosis; immunoproliferative disease; myphadenopathy; Alzheimer's disease; anagioimmunoproliferative lymphadenopathy; Alzheimer's disease; anagioimmunoproliferative lymphadenopathy; munoblastive lymphadenopathy; homestive lymphadenopathy; homes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecular antigen array; vaccine; antigen; antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bacteriophage Q-beta Al coat protein mutant Qbeta-259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABG80632 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein;
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1; Mismatches
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No. 4.5e-71;
                                                    immunoblastive lymphadenopathy;
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Query Match
Best Local Similarity
Matches 131; Conserv

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Conservative

99.5%; 99.2%;

Score 661; DB 5; Pred. No. 4.5e-71;

Length 132;

0

Gaps

0,

Mismatches

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CC molecular scaffold comprising: (i) a core particle selected from: (1) a core particle of a non-natural origin; and (2) a core particle of natural corigin; and (ii) an organiser comprising at least one first attachment correct the organiser is connected to the core particle by at least one covalent bond; (b) an antigen or antigenic determinant with at least cone second attachment site, where the antigen or antigenic determinant is concurring with the antigen or antigenic determinant; and (ii) an cocurring with the antigen or antigenic determinant; and (iii) an cocurring with the antigen or antigenic determinant; and (iii) an cocurring with the antigen or antigenic determinant; and this second determinant, where the second attachment site is capable of association concurring with the antigen or antigenic determinant; and this second interact through at least one non-peptide bond to the first attachment site; and where the antigen or antigenic determinant and the scaffold interact concurring through the association to form an ordered and repetitive antigen array. The composition is used in immunisation and as a vaccine for diseases such as influenza, graft corporated and repetitive antigen array. The composition is used in course in the second stream and the scaffold interact corporate and interact producting and interact producting array. The composition is used in course in the second and trapetities antigen array. The composition is used in course in the second and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and interact producting and i
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Renner WA,
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04-MAX-2001; 2001US-0288549P.
05-CCT-2001; 2001US-032699P.
07-NOV-2001; 2001US-0331045P.
                                 virus like particle or bacterial protein (the scaffold protein) present sequence is bacterial protein or peptide which is coupl modified antigen to form the molecular antigen array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-636514/68.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 18; Page 146;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JAN-2002;
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Bachmann
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1, Tissot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   composition
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ţA,
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Sebbel
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                                                                            response against an antigen or a virus-like particle in an animal. (C) comprises a virus-like particle (VLP) bound to at least one antigen, or a VLP capable of being recognised by the immune system of the animal. Also described: (1) enhancing an immune response against an antigen or a VLP in an animal comprising introducing (C) into the animal; (2) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; (3) immunising or treating an animal comprising administering the vaccine to the animal, or priming or boosting a T cell response in the animal by administering the vaccine; and (4) enhancing anti-viral protection in an animal comprising introducing (C) into the animal. (C) has cytostatic, virucide, antibacterial, antiparasitic, fungicide, antitheroid, antiflammatory, antitheroid, antiflammatory, antitheroid, antiflammatory.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or a virus --
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antigen presenting cell; APC; immune response; virus like particle; VLP; cytostatic; virucide; antibacterial; antiparasiic; fungicide; antiallergic; immunosuppressive; antialdictive; antiinflammatory; antithyroid; antidiabetic; neuroprotective; nootropic; osteopathic; antirheumatic; antiarthritic; vaccine; immunisation; infectious disease; anti-viral protection; tumour; allergy; drug addiction; Crohn's disease; graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis Alzheimer's disease; osteoporosis; rheumatoid arthritis; inflammatory autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A composition, useful for enhancing or a virus-like particle, enhancing or immunizing or treating tumors or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteriophage Qbeta.
                                  antithyroid, antidiabetic, neuroprotective, nootropic, osteopathic, antirheumatic and antiarthritic activities. (C) or the vaccines can
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a composition (C) for enhancing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 182; 243pp; English
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anti-viral protection in an animal,
infectious diseases, e.g. viral
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      an antigen or
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                                                                                                                                                                                                                                                                                                                                                                                                                               05-OCT-2001; 2001US-0326998P.

07-NOV-2001; 2001US-0331045P.

18-JAN-2002; 2002US-00050902.

21-JAN-2002; 2002WO-1B000166.

19-JUL-2002; 2002US-0396637P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cardiant, nephrotropic, ophthalmological; immunostimulant; vaccine, angiotensin peptide moiety carrier conjugate; angiotensin peptide; renin-activated angiotensin system; hypertension; stroke, infarction congestive heart failure; kidney failure; retinal haemorrhage; mutan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage Qbeta mutant coat protein
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New angiotensin peptide moiety carrier conjugate comprising a carrier and an angiotensin peptide moiety, useful for treating or preventing a disorder associated with renin-activated angiotensin, e.g. hypertension

2003-430264/40

Claim 20;

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97pp;

English.

invention

describes an angiotensin peptide moiety carrier conjugate

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Search completed: January 4, 2005, 09:11:02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    comprising: (a) a carrier with at least one first attachment site; and (b) at least one angiotensin peptide moiety with at least one second attachment site. The angiotensin peptide conjugate and compositions comprising them are useful for immunising an animal against an including peptide, and for treating or preventing a physical disorder angiotensin peptide, and for treating or preventing a physical disorder associated with renin-activated angiotensin system such as hypertension, stroke, infarction, congestive heart failure, kidney failure, and retinal haemorrhage. The conjugate is also useful for inducing immune responses, including producing antibodies. This is the amino acid sequence of a mutant bacteriophage Obeta coat protein used in the preparation of the vaccine conjugates of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 132 AA;
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US-08-232-144-4
US-08-232-144-4
US-09-200-673-15
US-10-1013-846-4
US-09-200-673-15
US-09-708-392-9
PCT-US93-05039-3
DCT-US93-05039-3
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DCT-US93-1437-3
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US-09-270-767-44620
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US-08-725-459B-43
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US-08-725-459B-42
US-09-248-796A-20249
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RESULT 2
US-09-948-722-2
; Sequence 2, Application US/09948722
; Patent NO. 6776993
; GENERAL INFORMATION:
; APPLICANT: Kaufmann, Stefan H. E.
; APPLICANT: Hess, Jurgen
; TITLE OF INVENTION: Tuberculosis Vaccine

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CURRENT FILING DATE: 2002-04-08
PRIOR APPLICATION NUMBER: US 09/485,717
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: PCT/EP98/05109
PRIOR FILING DATE: 1998-08-12
PRIOR APPLICATION NUMBER: EP 97/114,614.7
PRIOR FILING DATE: 1997-08-23
                                                                                                                                                                                                                                                                                                                                                                 US-09-328-352-4249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: GARY L. Breton et al.
APPLICANT: GARY L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
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SOFTWARE: PatentIn version 3.1
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Best Local
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CURRENT FILING DATE: 1999-06-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: recombinant nucleic acid OTHER INFORMATION: molecule comprising a domain of Mycobacterium and a phagolysomal OTHER INFORMATION: escape domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                                                                                                  109 SLINFTKDVRGGTRLNVNVPRNALISFAVNPVDGTTPI----IAKPNQRLTIRDLLAPG 163
                                                                                          164 RTGSN---AIAYLRETGFTNNAALVPENTAKPYSEITF--
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                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                         Indels
                                                                                          ----EEVMESVKTIAHMLK 213
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CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 23637
ENERGYH: 916
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT ; ORGANISM: Pseudomonas aeruginosa US-09-252-991A-23637
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 23637, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/09045186 Patent No. 6087154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local 9
                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/045,186
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Gaylo, Paul J.
REGISTRATION NUMBER: 36,808
REGISTRATION NUMBER: 36,808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                             TELEFAX: (317) 276-3861
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 3
                                                   SEQUENCE CHARACTERISTICS: LENGTH: 383 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: RHESUS NEUROPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (317) 276-0756
MOLECULE TYPE:
                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                     IOPOLOGY:
                                                                                                                                      TELEPHONE:
                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                     linear
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NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11.2%; Score 74.5; Di 23.7%; Pred. No. 5.3;
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WS-08-555-268A-15
US-08-555-268A-15
; Sequence 15, Application US/08555268A
; Patent No. 5958709
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Best Local Similarity 25.9
28; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT AFFLICATION NUMBER: US/08/234,
APPLICATION NUMBER: US/08/234,
FILING DATE: 26-MAY-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ERNST, Barbara G
NAME: ERNST, Barbara 30,377
                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SELBIE, Lisa
APPLICANT: HERZOG, Herbert
APPLICANT: SHINE, John
TITLE OF INVENTION: Human Neuropeptide Y-Y1 Receptor
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rottwell, Figg, Ernst & Kurz
STREET: 555 13th St, N.W., Suite 701-East
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.24 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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                                                                                                                                                                                                        16 ROTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
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                                                                                                                                      68 IQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAAL 115
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                                                                                                                                                                        RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN---
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                                                                                                    VTLDAYKDKYVCFDOFPSDSHRLSYTTLLLVL 218
                                                                                                                                                                                                                                        ; Score 71.5; D; Pred. No. 3.2; 11; Mismatches
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Pred. No. 3.
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                                                                                                                                                                                                                                                                          DB 1;
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                                                                                                                                                                                                                                            34; Indels 35;
                                                                                                                                                                                                                                                                         Length 384;
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Best Local (
                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                      Patent No.
                                                                                                                                                                                                                                                                                                                    Sequence 15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                 APPLICANT: Walker, Mary W.
APPLICANT: Branchek, Theresa
APPLICANT: Exanchek, Theresa
TITLE OF INVENTION: Methods of Modifying Feeding Behavior, Compounds Useful
TITLE OF INVENTION: in Such Methods, and DNA Encoding A Hypothalamic
TITLE OF INVENTION: Atypical Neuropeptide Y/Peptide YY Receptor (Y5)
FILE REFERENCE: 46166-BZ/JPW
CURRENT FILLING DATE: 1998-11-25
EARLIER APPLICATION NUMBER: US/09/200,673A
CURRENT FILLING DATE: 1998-11-25
EARLIER APPLICATION NUMBER: 08/566,096
ERRLIER FILLING DATE: 1995-12-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 391-052
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bard,
                                                                                                                                                                                                                                     APPLICANT: Gerald, Christophe P.G. APPLICANT: Weinshank, Richard L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 44743-Z/JPW/MAT
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
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APPLICANT: Weinshank, Richard L.
TITLE OF INVENTION: DNA ENCODING A HUMAN NEUROPEPTIDE
TITLE OF INVENTION: Y/PEPTIDE YY/PANCREATIC POLYPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ELECURATE (212)
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CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    138 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN------
FILING DATE: 1994-12-02
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                    APPLICATION NUMBER: 08/349,025
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1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             384 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10.8%; Score 71.5; DB 2; 25.9%; Pred. No. 3.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US/08/555,268A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11; Mismatches
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RESULT 10
US-09-708-392-9
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Best Local Similarity
Matches 28; Conserve
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APPLICANT: Bakthavatchalam, R
APPLICANT: Blum, Charles A
APPLICANT: Blum, Charles A
                                  Sequence 9, Application US/09708392
Patent No. 6734186
GENERAL INFORMATION:
APPLICANT: Pfizer Inc
APPLICANT: Pfizer Limited
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: NOO. 2001
CURRENT APPLICATION NUMBER: US/10/013,846
CURRENT FILING DATE: 2001-12-11
PRIOR APPLICATION NUMBER: US 60/254,990
PRIOR FILING DATE: 2000-12-12
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 15
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TYPE: PRT
ORGANISM: homosapiens
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ORGANISM: Homo sapiens
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Local Similarity 25.9%;
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5. 6566367
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Wayman,
Maw, G
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Elliott, Richard L
Hammond, Marlys
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Hutchison, Alan
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Pred. No. 3.2;
1; Mismatches 34;
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SEQ ID NO 9
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PRIOR APPLICATION NUMBER: US (
PRIOR FILING DATE: 2000-07-27
PRIOR FILING DATE: 2000-07-27
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PRIOR APPLICATION NUMBER: GB 0
PRIOR FILING DATE: 2000-07-05
PRIOR APPLICATION NUMBER: GB 0
PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRI
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PRIOR APPLICATION NUMBER:
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PRIOR FILING DATE: 2000-02-18
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CURRENT APPLICATION NUMBER: US/09/708,392
CURRENT FILING DATE: 2000-11-08
                              COMPUTER: Macintosh
OPERATING SYSTEM: MS-DOS
SOFTWARE: Microsoft Word 4.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/0
FILING DATE: 19930527
CLASSIFICATION:
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CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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APPLICATION NUMBER: US 60/175,161
TITLING DATE: 2000-01-07
TITLING DATE: 2000-01-07
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APPLICATION NUMBER: US 60/217,479
FILING DATE: 2000-07-11
APPLICATION NUMBER: US 60/221,014
FILING DATE: 2000-07-27
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                                                                                                                                                                                                                                                                                      floppy disk
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REGISTRATION NUMBER:

26,824

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RESULT 12
US-08-817-869-3
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                           TELEFAX: (732) 594-4720 INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           equence 3, Applic
                                                                                                                                                           REGISTRATION NUMBER: 25,147
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPANDED: PC-DOS/MS-DOS
COMPANDED: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: STRADER, CATHERINE D.
APPLICANT: CASCIERI, MARCARET A.
APPLICANT: MACNEIL, DOGLAS J.
TITLE OF INVENTION: MODIFIED NEUROPEPTIDE Y RECEPTORS
                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/335,017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: CR
TELECOMMUNICATION INFORMATION:
                                                                                            SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                 FILING DATE: 07-NOV-1994
ATTORNEY/AGENT INFORMATION:
NAME: NICHOLSON, WILLIAM H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: RAHWAY
STATE: NEW JERSEY
                   TOPOLOGY:
                                STRANDEDNESS:
                                                       TYPE: amino acid
                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           187 --
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                138 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEDFQN--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16 RQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IONPTACTANGSCDPSVTROKYADVTFSFTQYSTDEERAFVRTELAAL 115
TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/08817869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             126 EAST LINCOLN AVENUE
                                                                       411 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WILLIAM H. NICHOLSON
                                                                                                                                                (732) 594-1348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (203) 268-1951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10.8%; Score 71.5; I
25.9%; Pred. No. 3.2;
Live 11; Mismatches
                                                                                                                                                                                                                                                                                                                                                    US/08/817,869
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                                                                                                                                                                                                                                                                                                                                                                                          Version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34;
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Query Match

10.8%;

Score 71.5;

DB 3;

Length 411;

RESULT 14 US-09-270-767-44620

; Sequence 44620, Application US/09270767
; Patent No. 6703491

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                                                                                                                                Matches
                                                                                                                                              Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 34,087
REFERENCE/DOCKET NUMBER: 19:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-3462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT/US95/14377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: STRADER, CATHERINE U.
APPLICANT: CASCIERI, MARGARET A.
APPLICANT: MACCHEIL, DOUGLAG, NOTERING MET
                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 07-NOV-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16 ROTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: US
ZIP: 07065-0900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                ENGTH:
                                                              141 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             141 RHQLIINPRGWRPNNRHAYVGIAVIWVLAVASSLPFLIYQ--VMTDEPFQN------
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                             68 IONPTACTANGSCDPSVTROXYADVTFSFTQYSTDEERAFVRTELAAL 115
                                                                                             16 RQTLVLNPRGVNPTN-----GVA---SLSQAGAVPALEKRVTVSVSQPSRNRKNYKVQVK 67
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                                                                                                                                28;
                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                   APPOLLINA, MARY
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                                                                                                                                                                                                                                                                              411 amino acids
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                                                                                                                                Conservative
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                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                            peptide
                                                                                                                                                                                                                                               single
                                                                                                                                                                                                                                                                                                                                594-4720
                                                                                                                                               10.8%;
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VILDAYKOKYVCFDQFPSDSHRLSYTTLLLVL
                                                                                                                                11; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Pred. No. 3.6; 
11; Mismatches
                                                                                                                                              Score 71.5; DI
Pred. No. 3.6;
                                                                                                                                                                                                                                                                                                                                                                                      19339Y PCT
                                                                                                                                                               DB 5;
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                                                                                                                                                             Length 411;
                                                                                                                                Indels
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                                                                                                                                35;
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                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 44620
LENGTH: 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/08090148
Patent No. 5534257
GENERAL INFORMATION:
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Best Local Similarity
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                                                                               TELEFAX: (212) 940-640 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Drosophila melanogaster
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10022-2585

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5", 1.44Mb

COMPUTER: IBM PS2-486

COMPUTER: MS-DOS

OPERATING SYSTEM: MS-DOS
                                                            SEQUENCE CHARACTERISTICS:
                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 940-8636
TELEFAX: (212) 940-64064
FORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: Nisserbaum, Israel
REGISTRATION NUMBER: 27,1
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9101550.3
FILING DATE: 01/24/91
APPLICATION NUMBER: PCT/GB92/00124
FILING DATE: 01/22/92
                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/090,148
FILING DATE: 08/11/93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                  TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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IOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        315 AXXVVKYFSCTSSATRHSTRCISSYVSQSLGPSPLLGXADXKDFVRTLXPLATFISSPNL 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          258 TIAVN---ISPSPSPTPQPIVEPVEQVENSVTVTVASPEVPPVAXARNDRXRQTRSTIKN 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18 TLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQP-----SRNRKNYKVQVKIQN 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ΝY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3: Rosenman & Colin
575 Madison Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mastico, Robert Allan
Stockley, Peter George
Talbot, Simon John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antigen-Presenting Capsid with Fusion MS2-Coat Protein
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24.2%; Pred. No. 6.6;
tive 20; Mismatches
                                                                                                                                                                                                                                                                                       PCT/GB92/00124
                                                                                                                                                                                               27,582
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Search completed: January 4, 2005, 09:35:45
Job time: 10.3622 secs
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                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 24; Conserv
                                                                                                                                                                                                                                                                                                               VOLUME: 22-88
PAGES: 82-88
                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: Sequence.

PUBLICATION INFORMATION:

AUTHORS: Min Jou, W.; Haegeman, G.;

AUTHORS: Ysebaert, M.; Fiers, W.

TITLE: Nucleotide sequence of the

TITLE: gene coding for the

TITLE: bacteriophage MS2 coat protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: PROTEIN ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                               JOURNAL: Nat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY:
                                                                              77 LPVAAWRSYLNMELTIPIFATNSDCELIVKAMQGLL 112
                                                                                                                                                      24 NFANGVAEWISSNSRSQA-----YKVTCSVRQSSAQNRKYTIKVEVPKVATQTVGGVE 76
                                                                                                                                                                                             27 NPTNGVA-----SLSQAGAVPALEKRYTVSVSQPSRNRKNYKVQVKIQNPTACTANGSC
                                                                                                                                                                                                                               24; Conservative
                                                                                                                DPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALL 116
                                                                                                                                                                                                                                                                                                                                                                                        Nature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location 1 through 129 below represents entire MS2 coat protein in the published
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Coat protein from MS2-RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bacteriophage
                                                                                                                                                                                                                               10.5%; Score 69.5; DB 1; Length 129; 25.0%; Pred. No. 1; tive 16; Mismatches 43; Indels 1
                                                                                                                                                                                                                                     13;
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Run on:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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13:
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cgn2_6/ptodata/1/pubpaa/PCT N
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//Ggn2_6/ptodata/1/pubpaa/US10_RUB_PUB_pep:*
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SUMMARIES

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100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0	Query Match
132 132 132 132 132 132 132 132 132	% Query Match Length
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US-10-243-739-27 US-10-244-065-27 US-10-289-454-27 US-10-050-902-258 US-10-050-898-258 US-10-346-190-27 US-10-346-191-18 US-10-465-811-18 US-10-289-456-27 US-10-622-104-21 US-10-622-124-21 US-10-622-087-21 US-10-243-739-26 US-10-244-065-26	ID
Sequence 27, Appl Sequence 27, Appl Sequence 27, Appl Sequence 28, App Sequence 258, App Sequence 27, Appl Sequence 27, Appl Sequence 27, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl	Description

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US-10-243-739-10	-10-622-087-	-10-622	-10-622-064-	-10-289-456-	-10-465-811-	-10-346-190-	0-050-898-25	-10-050-902-	-10-289-454-2	-10-244-065-2	-739-	-10-622	-10-622-124-	-10-622-064-	-289-456-	0 - 465 - 8	-346-190-	-10-050-898-	-10-050-902-2	-10-289-454-	-10-244-065-	-10-243-739-	-10-622-087-	-10-622-124-2	22-064-	0-289-456-	0-465-811-	0-2	98-2	US-10-050-902-259	
Sequence 10, Appl	e 17	17	Sequence 6, Appli	e 23	e 14,	e 23	e 25	e 255,	e 23,	e 23,	e 23,	19,	e 19	e 8,	ø	e 16	25	e 25	e 25	e 25	e 25	25	20	20	e 9,	æ	e 17	e 26	equence 25	Ü	

ALIGNMENTS

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; TYPE: PRT
; ORGANISM: Bacteriophage Q-beta
US-10-243-739-27
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CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 27, Application US/10243739
Publication No. US20030091593Al
GENERAL INFORMATION:
APPLICANT: Bachmann, Martin F.
APPLICANT: Storni, Tazio
APPLICANT: Lechner, Franziska
APPLICANT: Lechner, Franziska
APPLICANT: Lechner, Franziska
TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of
TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
FILE REFERENCE: 1700.021001
                                                                                                                                                                                                                                                  Query Match 100.0%; Score 664; DB 14; Best Local Similarity 100.0%; Pred. No. 1.6e-66; Matches 132; Conservative 0; Mismatches 0;
121 LIDAIDQLNPAY 132
                                                  61
                                                                                        61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
                                                                                                                                                                             1 ARLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKKVTVSVSQPSRNRK 60
                                                  NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
                                                                                                                                                        ARLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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CURRENT APPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR PPLICATION NUMBER: PCT/IB02/00166
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
                                                                                                                                                                                                                                                                                          Sequence 27, Application US/10289454
Publication No. US20030157479A1
GENERAL INFORMATION:
APPLICANT: Bachmann, Martin
APPLICANT: Jennings, Gary
APPLICANT: Sonderegger, Ivo
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Best Local Similarity 100.0%;
Matches 132; Conservative (
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Publication No. US20030099668A1
                                                                                                                                                                                                                                             APPLICANT: Jennings, Gary
APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases
FILE REFERENCE: 1700.0360001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/244,065 CURRENT FILING DATE: 2002-09-16 PRIOR APPLICATION NUMBER: 60/374,145 PRIOR FILING DATE: 2002-04-22 PRIOR APPLICATION NUMBER: 60/318,994 PRIOR FILING DATE: 2001-09-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Partic TITLE OF INVENTION: Method of Preparation and Use FILE REFERENCE: 1700.0220001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 73
SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61
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Lipowsky, Gerad
Pumpens, Paul
Cielens, Indulis
Renhofa, Regina
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Pred. No. 1.6e-66;
); Mismatches '0;
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CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/321,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. SEQ ID NO 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 386
SOFTWARE: PatentIn version 3.2
SEQ ID NO 27
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                                                                                                                                                                                                                                                           Matches
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Best Local Similarity
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APPLICANT: Bachma
APPLICANT: Tissot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Piossek, Christine TITLE OF INVENTION: Molecular Antigen Array FILE REFERENCE: 1700.0190004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                             LENGTH: 132
TYPE: PRT
ORGANISM: Qb 259
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                                     121 LIDAIDQLNPAY 132
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                                                                                    61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
                                                                                                             61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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LIDAIDQLNPAY 132
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Sebbel, Peter
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Maurer, Patrick
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100.0%; Pred. No. 1.6e-66;
htive 0; Mismatches 0;
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US-10-346-190-27
Sequence 27, Application US/10346190
Publication No. US20030219459A1
GENERAL INFORMATION:
APPLICANT: Bachmann, Martin
APPLICANT: Maurer, Patrica
APPLICANT: Melicani, Erica
APPLICANT: Renner, Wolfgang A.
TITLE OF INVENTION: Prion Protein Carrier-Conjugates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 5
US-10-050-898-258
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; ORGANISM: Qb 259
US-10-050-898-258
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CURRENT APPLICATION NUMBER: US/10/050,898
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION DATE: 2001-11-07
PRIOR FILING DATE: 2001-11-07
FILE REFERENCE: 1700.0290003
; CURRENT APPLICATION NUMBER: US/10/346,190
; CURRENT FILING DATE: 2003-01-17
; PRIOR APPLICATION NUMBER: 60/396,590
PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/393,725
; PRIOR FILING DATE: 2002-07-08
; PRIOR APPLICATION NUMBER: 60/389,898
; PRIOR APPLICATION NUMBER: 60/389,898
; PRIOR APPLICATION NUMBER: 60/389,898
; PRIOR FILING DATE: 2002-06-20
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LENGTH: 132
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Maurer, Patrick
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Sebbel, Peter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Piossek, Christine
Ortmann, Rainer
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; TYPE: PRT
; ORGANISM: Qb
US-10-346-190-27
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US-10-465-811-18
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Publication No. US20040005338A1
GENERAL INFORMATION:
APPLICANT: BACHMANN, MARTIN F
APPLICANT: RENNER, WOLFGANG A
TITLE OF INVENTION: METHOD OF PREPARATION AND USE AS ADJUVANTS:
TITLE OF INVENTION: METHOD OF PREPARATION AND USE
FILE REFERENCE: 1700.029004
CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: US/10/465,811
CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: US 60/389,898
PRIOR FILING DATE: 2002-06-20
NUMBER OF SECTION OF NOC. 131
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Best Local S
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SOFTWARE: PatentIn version 3.1
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PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: 10/050,902
PRIOR FILING DATE: 2002-01-18
                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 131
                                                                                                                                                                                                                                                                                                                            TYPE: PRT
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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121
                        121 LIDAIDQLNPAY 132
                                                                                                                                                                                                                     132; Conservative
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                                                                      61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 ARLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60
                                                                                                                                                                                 1 ARLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60
                                                                                                                                                                                                                  h 100.0%; Score 664; DB 15; Similarity 100.0%; Pred. No. 1.6e-66; 32; Conservative 0; Mismatches 0;
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                                                                                               NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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LIDAIDQLNPAY 132
                                                                                                                                               ARLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60
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US-10-289-456-27; Sequence 27, Application US/10289456

RESULT 8

Publication No. US20040033211A1

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PRIOR APPLICATION NUMBER: US 60/396,575
PRIOR FILING DATE: 2002-07-18
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PatentIn version 3.2
SEQ ID NO 10
LENGTH: 132
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 9
US-10-622-064-10
; Sequence 10, Application US/10622064
; Publication No. US20040059094A1
; GENERAL INFORMATION:
                                                                                                                                                     ; OTHER INFORMATION: Bacteriophage Q-beta 259 mutant
US-10-622-064-10
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CURRENT FILING DATE: 2002-11-07
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR PILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2002-07-19
PRIOR PILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 170
                                                                                      Query Match
Best Local Similarity
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                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Maurer, Patrik F
TITLE OF INVENTION: Hapten-Carrier Conjugates and Uses Thereof
FILE REFERENCE: 1700.0300001
CURRENT APPLICATION NUMBER: US/10/622,064
CURRENT FILING DATE: 2003-07-18
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bachmann, Martin F APPLICANT: Maurer, Patrik F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Bachmann, Martin
APPLICANT: Maurer, Patrick
APPLICANT: Spohn, Gunther
TITLE OF INVENTION: Antigen Arrays for Treatment of Bone Disease
FILE REFERENCE: 1700.0330001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Q-beta 259 mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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ARLETYTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRYTYSVSQPSRNRK 60
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                                                                   Conservative
                                                                                      100.0%;
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                                                                                    Score 664; DB 15;
Pred. No. 1.6e-66;
                                                                   Mismatches
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                                                                                                         DB 15;
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US-10-622-124-21
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PRIOR FILING DATE: 2002-07-19
NUMBER OF SEQ ID NOS: 146
SOFTWARE: PatentIn version 3.2
SEQ ID NO 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21, Application US/10622124 Publication No. US20040076645A1 GENERAL INFORMATION:
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Best Local 9
PRIOR APPLICATION NUMBER: US 60/396,639
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: US 60/470,432
PRIOR FILING DATE: 2003-05-15
NUMBER OF SEQ ID NOS: 93
SOFTWARE: PatentIn version 3.2
SEQ ID NO 21
                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/622,087
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                      TITLE OF INVENTION: Amyloid Beta 1-6 Antigen Arrays FILS REFERENCE: 1700.0350002
                                                                                                                                                                                                                                 APPLICANT:
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APPLICANT: Tissot, Alain
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CURRENT FILING DATE: 2003-07-18
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TITLE OF INVENTION: Ghrelin-Carrier Conjugates
FILE REFERENCE: 1700.0340001
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APPLICANT: Fulurija, Alma
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 132
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100.0%; Pred. No. 1.6e-66;
tive 0; Mismatches 0;
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US-10-244-065-26
; Sequence 26, Application US/10244065
; Publication No. US20030099668A1
; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Bacteriophage Q-beta 259 mutant
US-10-622-087-21
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; TYPE: PRT
; ORGANIZM: Bacteriophage Q-beta
US-10-243-739-26
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APPLICANT: Storni, Tazio
APPLICANT: Storni, Tazio
APPLICANT: Lechner, Franziska
TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of
TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles
FILE REFERENCE: 1700.0210001
CURRENT APPLICATION NUMBER: US/10/243,739
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/318,967
PRIOR PRILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE: Patentin version 3.1
SEQ ID NO 26
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                  APPLICANT: Bachmann,
APPLICANT: Storni,
APPLICANT: Maurer,
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nes 132; Conser
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Storni,
Maurer,
Tissot,
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                                       Tazio
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                                                          Martin F.
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Pred. No. 3.5e-66;
1; Mismatches 0;
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APPLICANT: Sonderegger, Ivo
TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eos:
FILE REFERENCE: 1700.0360001
CURRENT APPLICATION NUMBER: US/10/289,454
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 60/396,636
PRIOR FILING DATE: 2002-07-19
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2002-01-11-07
PRIOR PELLING DATE: 2001-11-07
NUMBER: OF SEQ ID NOS: 386
PRIOR FILING DATE: 2001-11-07
NUMBER: DETAILS 2001-11-07
NUMBER: DETAILS 2001-11-07
                                                                                    ; SEQ ID NO 26
; LENGTH: 132
; TYPE: PRT
; ORGANISM: Bacteriophage Q-beta-251
US-10-289-454-26
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APPLICANT: Renhofa, Regina
TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Particle
TITLE OF INVENTION: Method of Preparation and Use
FILE REFERENCE: 1700.0220001
CURRENT APPLICATION UNMEER: US/10/244,065
CURRENT FILING DATE: 2002-09-16
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/374,145
PRIOR APPLICATION NUMBER: 60/318,994
PRIOR FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 73
SOFTWARE OF SEQ ID NOS: 73
SOFTWARE OF SEC ID NOS: 73
               Query Match
Best Local Similarity
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    Matches
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Best Local Similarity
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APPLICANT:
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APPLICANT: Jennings, Gary
                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.2
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ORGANISM: Bacteriophage Q-beta
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Lipowsky, Gerad
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    Conservative
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                    99.5%;
99.2%;
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Score 661; DB 14;
Pred. No. 3.5e-66;
1; Mismatches 0;
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Pred. No. 3.5e-66;
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1 ARLETVTLGNIGKDGRQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK 60

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APPLICANT: Renner, Woligany A.
APPLICANT: Bachmann, Martin
APPLICANT: Tissot, Alain
APPLICANT: Lechner, Patrick
APPLICANT: Lechner, Franziska
APPLICANT: Lechner, Franziska
APPLICANT: Sebbel, Peter
APPLICANT: Piossek, Christine
FILE REFERENCE: 1700.0190004
CURRENT APPLICATION NUMBER: US/10/050,902
CURRENT APPLICATION NUMBER: US 60/262,379
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-01-0
PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-10-05
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 350
SOFTWARE: Patenetin Ver. 2.1
SEQ ID NO 259
LENGTH: 132
TYPE: PRT
ORGANISM: Qb 251
US-10-050-902-259
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US-10-050-902-259
; Sequence 259, Applicat
; Publication No. US2003
; DENERAL INFORMATION:
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Search completed: January 4, 2005, 09:41:23 Job time : 32.7351 secs
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                                                                                                                                                        61 NYKVQVKIQNPTACTANGSCDPSVTRQKYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
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Maximum Match 100%
Listing first 45 s
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Maximum DB
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Perfect score:
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  Pred. No.
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116
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seq length: 2000000000
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Match
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664
1 ARLETVTLGNIGKDO
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Gapop 10.0 ,
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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 ARLETVTLGNIGKDGRQTLV.....AALLASPLLIDAIDQLNPAY 132
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(c) 1993 - 2005
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          VCBPQB

$01964

VCBPP1

$22340

T00646

$224231

A41505

A41505

A41166

T31666

T31666

T3165

T3165

T3165

T3165

T3165

T3165

T3165

T3165

T3165

T3167

T346324

VCBPFA

VCBPFA

VCBPFA

VCBPFA

VCBPFA

VCBPFA

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45	44	43	42	41	40	39	38	37	36	3 5	34	ω	32	31	30
69.5	70	70	70.5	71	71	71	71	71	71	71	71.5	72	72.5	72.5	72.5
10.5	10.5	10.5	10.6	10.7	10.7	10.7	10.7	10.7	10.7	10.7	10.8	10.8	10.9	10.9	10.9
130	425	243	836	1545	1545	1461	1461	463	399	248	384	376	129	129	129
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coat protein [vali	pectate lyase (EC	nucleocapsid prote	conserved hypothet	sulfonylurea recep	sulfonylurea recep	hypothetical prote	hypothetical prote	translation elonga	polymerase-associa	conserved hypothet	neuropeptide Y/pep	hypothetical prote	coat protein - pha	coat protein - pha	coat protein - pha

ALIGNMENTS

RESULT VCBPQB

coat protein - phage Q-beta
C;Species: phage Q-beta
C;Species: phage Q-beta
C;Date: 29-Jul-1981 #sequence_revision 24-Sep-1981 #text_change 09-Jul-2004
C;Accession: A92240; A92221; A92088; A04224
R;Escarmis, C.; Sastry, P.A.; Billeter, M.A.
J. Biol. Chem. 253, 8390-8399, 1978
A;Title: Determination of the first half of the coat protein cistron of bacteriophage Qbe
A;Reference number: A92240; MUID:79048469; PMID:361741
A;Accession: A92240

A; Molecule type: mRNA A; Residues: 1-80 <ESC>

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A;Cross-references: UNIPROT:P03615
R;Stoll, E; Wilson, K.J.; Reiser, J.; Weissmann, C.
U. Biol. Chem. 252, 990-993, 1977
A;Title: Revised amino acid sequence of Obeta coat protein between positions 1
A;Reference number: A92221; MUID:77118576; PMID:838709
A;Recession: A92221
A;Molecule type: protein
A;Residues: 1-60 <STO>
R;Maita, T.; Konigsberg, W.
U. Biol. Chem. 246, 5003-5024, 1971
A;Reference number: A92088; MUID:71288580; PMID:5570434
A;Recession: A92088; MUID:71288580; PMID:5570434
A;Recession: A92088
RESULT 2
$01964.
$01964.
C;Species: phage SP
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
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A;Residues: 1-21,'D',23-55,57-132 <MAI>
C;Superfamily: phage GA coat protein
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                                                                                                                                                                                                                                        61 NYKVQVKIQNPTACTANGSCDPSVTRQAYADVTFSFTQYSTDEERAFVRTELAALLASPL 120
                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                           AKLETYTLGNIGKDGKQTLVLNPRGVNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRK
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                                                                                                                                                           LIDAIDQLNPAY 132
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      98.2%; Score 652; DB 1;
97.7%; Pred. No. 1.6e-56;
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R;Haas, A.; Dumbsky, M.; Kreft, J.
Biochim. Biophys. Acta 1130, 81-84, 1992
A;Title: Listeriolysin genes: complete so
A;Reference number: S22340; MUID:92182010
                                                                                        seeligeriolysin -
C;Species: ".ie+~--
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A;Title: Analysis of the complete nucleotide sequence of the group IV RNA coliphage A;Reference number: S01963; MUID:88289362; PMID:3399390
A;Accession: S01964
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A; Residues: 1-11 cHA>
A; Cross-references: UNIPROT: P03616
C; Superfamily: phage GA coat prote:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: genomic RNA
A; Residues: 1-331 < INO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 3
                                                          Date: 22-Nov-1993; Accession: S22340
                                                                        ;Species: Listeria seeligeri
;Date: 22-Nov-1993 #sequence_revision 01-Dec-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Date: 30-Nov-1979 #sequence_revision 30-Nov-1979 #text_change;Accession: A04225
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                                                                                                                                                                                                                                                                                                                                                                 17.5%; ilarity 32.3%; Conservative
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                                                                                                      Listeria seeligeri
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s: complete sequence of ilo MUID:92182018; PMID:1543752
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Pred. No. 5.4e-44;
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              Listeria
                                                                          09-Jul-2004
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C;Accession: S24231

R;Rasmussen, O.F; Beck, T.; Olsen, J.E.; Dons, L.; Rossen, L.

Infect. Immun. 59, 3945-3951, 1991

A;Title: Listeria monocytogenes isolates can be classified into
A;Reference number: S24230; MUID:92040062; PMID:1937753
                                                                                                                                                                                                                                                       RESULT 6
S24231
A; Accession: S24231
A; Status: nucleic ac
A; Molecule type: DN
                                                                                                                                                               listeriolysin precursor - Listeria monocytogenes (strain 12067)
C;Species: Listeria monocytogenes
A;Variety: strain 12067
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Introns: 14/3
C; Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Federspiel, N.A.; Palm, C.J.; Conway, A.B.; Kurtz, D.B.; Conway, A.R.; Au, 
; Vysotskaia, V.S.; Yu, G.; Ecker, J.; Theologis, A.; Davis, R.W. 
submitted to the EMBL Data Library, February 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:P31830; EMBL:X60462; NID:g44144; A;Note: the authors translated the codon GCC for residue 287 C;Superfamily: dipeptide transport protein
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A;Status: preliminary
A;Molecule type: DNA
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A;Molecule type: DNA
A;Residues: 1-540 <FED>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein F3I6.9 - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 01-Feb-1999 #sequence_revision 01-Feb-1999 C;Accession: T00646
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A; Residues: 1-530 < HAA>
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                                                                                                                                                                                                                                                                                                                                           KPAAPMSKSPQGFATPRVYKPAPTKTSLSTSHSSLKKEK----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IGKDGRQTLVLNPRG----VNPTNGVASLSQAGAVPALEKRVTVSVSQPSRNRKN-YKVQV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                 acid sequence not shown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #sequence_revision 01-Feb-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              428/3; 448/2;
dehydrogenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12.9%; Score 85.5; DB 2; 26.1%; Pred. No. 1.8; tive 15; Mismatches 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                      translation
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                      not
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as Pro
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                                                                               major types according
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A;Accession: S12400
A;Molecule type: DNA
A;Residues: 483-493 <MIC>
A;Experimental source: str:
C;Genetics:
A;Gene: hlyA; lisA
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C;Species: Listeria monocytogenes
C;Date: 21-Oct-1992 #sequence revision 21-Oct-1992 #text_change 09-Jul-2004
C;Accession: A43505; 805306; A47606; S12400; A61079
R;Mengaud, J.; Vicente, M.F.; Chenevert, J.; Pereira, J.M.; Geoffroy, C.; Gicquel-Sanzey
Infect. Immun. 56, 766-772, 1988
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A;Residues: 1-529 <RAS>
A;Cross-references: UNIPOT:P13128; EMBL:X60035; NID:g44110; PIDN:CAA42639.1; PID:g44112
A;Experimental source: strain 12067, serotype 4b
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Title: Expression in Escherichia coli and sequence analysis of the listeriolysin O det A;Reference number: A43505; MUID:88153053; PMID:3126142
A;Accession: A43505
                                                                                                                                                       A; Molecule type: DNA
A; Residues: 413-460 «ME2»
A; Residues: 413-460 «ME2»
A; Cross-references: GB: M29171
A; Cross-references: GB: M29171
R; Michel, E.; Reich, K.A.; Favier, R.; Berche, P.; Cossart, P.
Mol. Microbiol. 4, 2167-2178, 1990
A; Title: Attenuated mutants of the intracellular bacterium Listeria
A; Reference number: S12400; MUID: 91211627; PMID: 1965218
                                                                                                                                                                                                                                                                                                                                                 R;Domann, E.; Chakraborty, T.
Nucleic Acids Res. 17, 6406, 1989
A;Title: Nucleotide sequence of the listeriolysin gene from a Listeria monocytogenes
A;Reference number: S05306; MUID:89366684; PMID:2505236
A;Accession: S05306
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A;Note: this sequence is derived from a strongly hemolytic strain, serotype 1/2c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-529 < MEN>
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;1-25/Domain: signal sequence #status predicted
;26-529/Product: listeriolysin #status predicted
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;Residues: 1-529 <DOM>
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                                                                                                                                                                                                                                                                                                                                       Status: preliminary
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                                                          strain LO28,
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22.7%;
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Pred. No. 2.
                                                       serotype 1/2c
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                                                                                                                                                                                    monocytogenes
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ivanolysin precursor - Listeria ivanovii (,Species: Listeria ivanovii C,Species: Listeria ivanovii C,Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_cha C,Accession: S22341; S36683 R,Haas, A.; Dumbsky, M.; Kreft, J. Biochim. Biophys. Acta 1130, 81-84, 1992 A,Title: Listeriolysin genes: complete sequence of ilo from A,Reference number: S22340; MUID:92182018; PMID:1543752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Superfamily: dipeptide transport protein C;Keywords: virulence factor F;1-25/Dmain: signal sequence #status predicted <SIG>F;26-529/Product: listeriolysin O #status predicted <M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Science 294, 849-852, 2001
A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; A; Authors: Kreft, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; ok, C.; Schlueter, T.; Simoes of Listeria species.
A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Accession: AC1100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-529 <GLA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Superfamily: dipeptide transport protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Gene: hly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; Species: Listeria monocytogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    listeriolysin O precursor [imported] - Listeria monocytogenes (strain EGD-e)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             h 12.7%; Score 84.5; D
Similarity 22.7%; Pred. No. 2.2;
30; Conservative 23; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                     TLSIDLPGMTNQDNKIVVK--NATKSNVNNAVNTLVERWNEKYAQAYPNVS---AKIDYD
                                                                                                                                                                                                                                                                                                                                                                                               TVSVSQPSRNRKNYKVQVKIQNPTACTANGSCDPSVTR-----
                                                                                                                                                                                                                                                                                                                                                                                                                                        KUGNEYIVVEKKKKSINQNNADIQVVNAISSLTYPGALVKANSELVENQPDVLPVKRDSL
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                                                                                                                                                                                                                                                                  DEMAYSESQLIA 219
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Dussurget, O.; Entian, K.D.; E
                                                                                                             #text_change 09-Jul-200
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                             Listeria ivanovii and
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Voss, H.; Wehland,
                                                                                                                                                                                                                                                                                                                                                                                                                                          152
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Fsihi, H.;
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A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-1502 < JOH>
A; Residues: 1-1502 < JOH>
A; Cross-references: EMBL: U20865; NID: g662330; PIDN: AAB67387.1; PID: g662331; GSPDB: GN0001
A; Experimental source: strain S288C (AB972)
A; Experimental source: strain S288C (AB972)
R; Pfeifer, K.; Kim, K.S.; Kogan, S.; Guarente, L.
Cell 56, 291-301, 1989
Cell 56, 291-301, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 10
RGBYH1
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A; Map position: 12R
C; Superfamily: regulatory protein HAP1;
                       A; Cross-references:
A: Map position: 12R
                                                                       A; Gene: SGD: HAP1; CYP1;
                                                                                                                  A; Note: the sequence is from
                                                                                                                                A;Molecule type: DNA
A;Residues: 1-62,'R',64-1305,'Y',1306-1470,'LVDFYRADFPIWE'
A;Cross-references: EMBL;X13793
                                                                                                                                                                                                          J. Mol. Biol. 204, 263-276, 1988
A;Title: CYPI (HAPI) regulator of oxygen-dependent gene expression in yeast.
A;Reference number: S15447; MUID:89125585; PMID:2851658
A;Accession: S15447
                                                                                                                                                                                                                                                                                             A;Residues: 1-144,'I',146-322,'R',324-454,'N',456-507,'M',509-586,'K',588-882,'N',884-95
A;Cross-references: EMBL:JO3152; NID:g171645; PIDN:AAA34662.1; PID:g171646
R;Creusot, F; Verdiere, J.; Gaisne, M.; Slonimski, P.P.
                                                                                                                                                                                                                                                                                                                                                                                          A;Title: Functional dissection and sequence of yeast HAP1 A;Reference number: A31312; MUID:89106221; PMID:2643482 A;Accession: A31312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CYC1/CYP3 transcription activator - yeast (Saccharomyces cerevisiae) N;Alternate names: protein L9672.1; protein YLR256w; regulatory prot
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C;Superfami
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A; Residues: 1-319, 'T
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Accession: S22341
A;Molecule type: DNA
A;Residues: 1-528 <HAS>
                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Accession: S59400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Description:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        submitted to the EMBL Data Library,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Accession:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Date: 30-Sep-1991 #sequence revision 23-Feb-1996 #text_change 12-Nov-1999; Accession: S59400; A31312; S15447; S05804; S15446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Species: Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;25-528/Product: ivanolysin #status predicted
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;Cross-references: EMBL:X60461
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                                             YP1; MIPS:YLR256w
SGD:S0004246; MII
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                                               MIPS:YLR256w
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Pred. No. 5
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  binuclear cluster homology
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    Overall

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hypothetical protein COS41.6 - sea squirt (Ciona intestinalis)
(;Species: Ciona intestinalis
C;Species: Ciona intestinalis
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 09-Jul-2004
C;Accession: T31660
R;Bird, A.P.; Clark, V.; Jones, S.J.; Leitgeb, S.; Dobson, R.; Tweedie, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 11
AI0186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; William Nature 413, 523-527, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probable iron-siderophore transport system, ATP-binding component YPO1533 [imported] C;Species: Yersinia pestis C;Species: Yersinia pestis C;Date: 02-Nov-2001 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              F;177-189/Region: glutamine-rich
F;245-445/Domain: heme binding Htatus predicted <HEM>
F;245-245/Domain: heme binding Htatus predicted <HEM>
F;299-304,323-328,347-352,373-378,389-394,415-420/Region:
F;1308-1481/Domain: activation element #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Keywords: DNA binding; heme binding; transcription regulation; F;1-148/Domain: DNA binding #status predicted <DNA> F;5-9-98/Domain: GAL4 zinc binuclear cluster homology <GAL4> F;64-84/Region: zinc finger CCCC motif
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C; Superfamily:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Title: Genome sequence of Yersinia pestis, the causative agent A;Reference number: AB0001; MUID:21470413; PMID:11586360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule
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                                                                                                                                                                                                                                                                                                        LLSQGASISETLTVFDLVSRGRYAHQSF-FHQWSTEDER-IVKAAISAVNLESLVQQRVS
                                                                                                                                                                                                                                                             QLN 129
                                                                                                                                                                                                                                                                                                                                                ACTANGSCDPSVT-----RQKYADVTFSFTQYSTDEERAFVRTELAALLASPLLIDAID 126
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                                                                                                                                                                                                                                                                                                                                                                                           KNGAFSVIIGPNGCGKSTLLRALSRS----LLPQNGSIRLDQ--QDIQHYKAKVFARQLS
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Library,
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Pred. No. 18;
    December
    1996
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R.M.; Davis, P.; Dougan, G.;
K.; Whitehead, S.; Barrell,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           63;
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Data

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C;Species: Saccharomyces cerevisiae C;Date: 10-Oct-1995 #sequence revision 10 C;Accession: S57527; S50279; S63024; S630 R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J submitted to the EMBL Data Library, June
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: T13845
R;Saunders, R.D.; Avides, M.C.; Howard, T.; Gonzalez, C.; Glover, D.M.
J. Cell Biol. 137, 881-890, 1997
A;Title: The Drosophila gene abnormal spindle encodes a microtubule-associated protein A;Reference number: Z17792; MUID:97296495; PMID:9151690
A;Accession: T13845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      microtubule-associated protein - fruit fly (Drosophila melanogaster) C_i; Species: Drosophila melanogaster
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                            A;Accession: S57537
A;Molecule type: DN
                                                                   A; Reference number: S57533
                                                                                                     Species: Saccharomyces cerevisiae
|Species: Saccharomyces cerevisiae
|Date: 10-Oct-1995 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004
|Accession: S57537; S50279; S63024; S63017; S65096
|Accession: S57537; S50279; S63024; S63017; S65096
|Accession: S57537; S50279; S63024; S63017; S65096
                                                                                                                                                                                                          KT1 protein - yeast (Saccharomyces cerevisiae),Alternate names: protein N2302, protein YNL08
Molecule type: DNA Residues: 1-830 <SOL>
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                                                                                                                                                                                                                                                                                                                                                                                                                                            ATPRKVRDTSIQPS--VKLYDSELYMQTCINPDPFAATTTIDPFLASTMYLD
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                                                                                                                                                                                                                                                                                                                                                      AVDRHQADFKKWLNALVSIPADLDA
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26.2%; Pred. No. 36;
/ative 17; Mismatches
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28.3%;
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Pred. No. 5.
                                                                                                                                                                                                            protein YNL085w
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                                             A; Cross-references:
                                                                        A;Residues: 1-136 <KUR>
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R;Goodner, B.; Hinkle, G.; Gattung, S.; Miller, N.; Blanchard, M.; Qurollo, A.; Liu, F.; Wollam, C.; Allinger, M.; Doughty, D.; Scott, C.; Lappas, C.; Science 294, 2323-2328, 2001
A;Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrob A;Reference number: A97359; MUID:21608551; PMID:11743194
A;Accession: C98221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Residues: 1-29,'G',31-808,'TMKTCLNYH' <VER>
A;Cross-references: EMBL:U09129; NID:g520475; PIDN:AAC49470.1; PID:g520476
R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M.
submitted to the Protein Sequence Database, April 1996
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Yeast 10, 1477-1479, 1994
Ä;Title: Sequence of MKT1, needed for propagation of M(2) satellite deRNA A;Reference number: S50279; MUID:95176705; PMID:7532890
                                                                                                                                                                                                                                                        hypothetical protein AGR_L_1428 [imported] - Agrobacterium tumefaciens (strain C;Species: Agrobacterium tumefaciens C;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 09-Jul-2004
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P;615-631/Domain: transmembrane #status predicted <TWM>
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C; Superfamily: Saccharomyces cerevisiae MKT1 protein
C; Keywords: transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast 12, 485-491, 1996
A,Title: The sequence of a 17 933 bp segment of Saccharomyces A;Reference number: S65092; MUID:96310628; PMID:8740422
A;Accession: S65096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-830 <SOF>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Experimental source: strain S288C
R;Soler-Mira, A.; Salz, J.B.; Ballesta, J.P.G.; Remacha, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Residues: 569-830 <POE>
A;Cross-references: EMBL:Z71361; MIPS:YNL085w
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A; Residues: 1-830 < SOW>
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A; Molecule type: DNA
                              A;Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: nucleic acid sequence not shown
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A;Experimental source: strain S288C
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UNIPROT: Q8U8F3; GB: AE007870; PIDN: AAK89293.1; PID: g15159127; GSPDB: GN

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Result
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Q9vc45 drosophila	Aar97352 listeria	Q6r6d0 listeria iv	Q6hga7 bacillus th	P91584 ciona intes	Aas61663 yersinia	Q8zfz0 yersinia pe	P12351 saccharomyc	Aar97343 listeria	Q6r6d9 listeria iv	P31831 listeria iv	Q8trf3 methanosarc	Q95ru8 drosophila	Q9v3k8 drosophila

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	ISS-PROT entry is copyright. It is produced through a collaboration the Swiss Institute of Bioinformatics and the EMBL outstation opean Bioinformatics Institute. There are no restrictions on its one-profit institutions as long as its content is in no way and this statement is not removed. Usage by and for commercial	OMS). dule M., Valegard K., Liljas L.; ophage Q-beta at 3.5-A resolution."; ll; binds to the phage RNA.	hem. 252:990-993(1977). 288580; PubMed=5570434; Konigaberg W.; acid sequence of the Q-beta coat	"; 'hem. 253:8390-8399(1978). TO 1-60. 118576; PubMed=838709; Wilson K.J., Reiser J., Weissmann C.; Wilson K.J., Reiser J., Weissmann C.;	1-80 FROM N.A. 8469; PubMed=361741; Sastry P.A., Billeter M.A.; on of the first half of the coat protein cistron of continuous conti	; linna D., Dislers A., Baumanis V., psid particles synthesized and self	(Rel. 01, Created) (Rel. 01, Last sequence update) (Rel. 45, Last annotation update) 1. 1. 1. 2. 2. 2. 2. 2. 2. 2. 2. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3. 3.	STANDARD; PRT; 132

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EMBL; AY099114; AAM33126.1; -. Coat protein.
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Pfam; PF01819; Levi_coat; 1.
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Q8LTE1;
Q9LTE1;
Q1-OCT-2002 (TrEMBLre1. 2
Q1-OCT-2002 (TrEMBLre1. 2
Q1-UT-2004 (TrEMBLre1. 2
Q5-UUL-2004 (TREMBLRE1. 2
A1 read-through protein (
Bacteriophage Q-beta;
AAA16663;
AAA16663;
02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; M99039; AAA16663.1; -.
HSSP; P03615; 1QBE.
GO; GO:0019928; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; IEA.
InterPro; IPR002703; Levi_coat.
InterPro; IPR00504; RNA_rec_mot.
Pfam; PF01819; Levi_coat; 1.
PROSITE; PS00030; RRM RNP 1; UNKNOWN 1.
SEQUENCE 329 AA; 36134 MW; E0F30215A8C197CB CRC64;
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PubMed=14667253;

Bacher J.M., Bull J.J., Ellington A.D.;

Bacher J.M., Bull J.J. Ellington A.D.;

"Evolution of phage with chemically ambiguous

BMC Evol. Biol. 3:24-24(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Viruses; ssRNA positive-strand viruses,
Allolevivirus.
NCBI_TaxID=12009;
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Q9TOR9;
01-MAY-2000
01-MAY-2000
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01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2003 (TrEMBLrel. 24,
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Kozlovska T.M., Cielens I., Dreilinna
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02-MAR-2004
                                                                                                                                                                                                                                             Beekwilder J., Nieuwenhuizen R., Poot R., van Duin J.; "Secondary structure model for the first three domains Control of A-protein synthesis."; J. Mol. Biol. 256:8-19(1996).
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Viruses; ssRNA positive-strand
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                                                                                                                     Beckwilder M.J., Nieuwenhuizen R., van Duin J.;
Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF055242; AAC14699.1; -.
HSSP; P03615; 1QBE.
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Pfam; PF01819; Levi_coat;
Coat protein.
SEQUENCE 133 AA; 14198
                                                         GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi_coat.
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MEDLINE=96190948; PubMed=8609616;
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Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
R EMBL; AF059242; AAC14700.1; -.
R GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; IEA.
R InterPro; IPR002703; Levi_coat.
R InterPro; IPR002703; Levi_coat.
R InterPro; IPR000504; RNA_rec_mot.
R InterPro; IPR001504; RNA_rec_mot.
R PROSITE; PS001030; RRM RNP 1; UNKNOWN 1.
SEQUENCE 329.AA; 35880 MW; 372B1DD5DDB52F15 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Beekwilder J. Nieuwenhuizen R., Pc "Secondary structure model for the Control of A-protein synthesis."; J. Mol. Biol. 256:8-19(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Secondary structure model RNA phage Q beta.";
J. Mol. Biol. 247:903-917()
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                                       LIDAIDQLNPAY 132
                                                                                     NYKVQVKIQNPTSCTASGTCDPSVTRSAYADVTFSFTQYSTDEERALVRTELKALLADPM
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larity 82.6%; Pred. No. 4.1e-46;
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Q9TOS0;
Q9TOS0;
01-MAY-2000
01-JUN-2000
01-JUN-2003
                                                               Al-protein.

Bacteriophage M11.

Viruses; ssRNA positive-strand
Allolevivirus.

NCBI_TaxID=74336;
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MEDLINE=95239761; PubMed=7723040;
Beekwilder M.J., Nieuwenhulzen R.,
"Secondary structure model for the
RNA phage Q beta.";
J. Mol. Biol. 247:903-917(1995).
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01-AUG-1998 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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Pfam; PF01819; Levi_coat; 1.
Coat protein.
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    SEQUENCE FROM N.A. MEDLINE=95239761;
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                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Ew the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content use by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
MEDLINE=88289362; PubMed=3399390;
Hirashima A., Hirose T., Inayama S.,
"Analysis of the complete nucleotide
coliphage SP.";
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InterPro; IPR000504; RNA rec mot.
Pfam; PF01819; Levi_coat; 1.
PROSITE; PS00030; RRM RNP 1; UNKNOWN 1
SEQUENCE 329 AA; 35893 MW; 3E33CD821EE625F4
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"Secondary structure model for the
Control of A.protein synthesis.",
J. Mol. Biol. 256:8-19(1996).
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                                                                                                entities requires a license agreement (Some send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatice Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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P09677;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic Acids Res. 16:6205-6221(1988).
-!- FUNCTION: Not yet known.
-!- MISCELLANEOUS: The readthrough protein
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01-MAR-1989 (Rel. 10, Last sequence update)
29-MAR-2004 (Rel. 43, Last annotation update)
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Pfam; PF01819; Levi_coat; 1.
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Pred. No. 3.1e-43;
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Readthrough protein A1.
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01-MAY-2000
Al-protein.
Enterobacteria phage NL95.
Viruses; ssRNA positive-strand viruses,
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GO; GO:0005198; F:structural molecule
InterPro; IPR002703; Levi_coat.
Pfam; PF01819; Levi_coat; 1.
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HSSP; P03615; 1QBE.
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3 (TrEMBLrel. 07,
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                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 74.8%; Score 496.5; DB 2; 75.0%; Pred. No. 3.9e-40; vative 11; Mismatches 21;
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ne first three domains
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                 no DNA stage; Leviviridae;
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Best Local :
                Fur. A04225; VCBpp1.

PIR; A04225; VCBpp1.

PIR; A04225; VCBpp1.

Levi_coat; 1

Tore;
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P03616;

21-JUL-1986

21-JUL-1986

05-JUL-2004
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GO; GO:0005198; F:structural molecule activity;
GO; GO:0005198; F:structural molecule activity;
InterPro; IPR002703; Levi coat.
InterPro; IPR000504; RNA rec_mot.
Pfam; PF01819; Levi coat; 1.
PROSITE; PS00030; RRM RNP 1; UNKNOWN 1.
SEQUENCE 330 AA; 36175 MW; 961E55F408334410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beekwilder M.J., Nieuwenhuizen R., "Secondary structure model for the RNA phage Q beta.";
J. Mol m:-"
                                                                                                                   SEQUENCE.
MEDLINE=79148387; PubMed=107028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Beckwilder M.J., Nieuwenhuizen R.,
Submitted (APR-1998) to the EMBL/G
EMBL, AF055243, AAC14704.1, -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        J. Mol. Biol.
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J. Mol. Biol.
            Pfam; PF01819; Levi c
                                                                                  bacteriophage PRR1."
                                                                                            The primary structure of the
                                                                                                                                                                   Viruses; ssRNA positive-strand viruses, unclassified Leviviridae.
                                                                                                                                                                                      Bacteriophage
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"Secondary structure model for the
Control of A-protein synthesis.";
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NCBI_TaxID=75725;
                                                         ur. J. Biochem. 94:375-386(1979).
                                                                                                                                                      _TaxID=12024;
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 131 AA;
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(Rel.
(Rel.
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 protein sequencing; RNA-binding 14535 MW; E7E639E1E50FC612 CRC6
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75.0%;
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EMBL/GenBank/DDBJ
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No. 1.1e-39;
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                                                                                             coat
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ne first three domains
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RESULT 15
TACY LISSE STANDARD;
ID TACY LISSE STANDARD;
AC P31830;
DT 01-JUL-1993 (Rel. 26, Create
DT 01-JUL-1993 (Rel. 26, Last &
DT 05-JUL-2004 (Rel. 44, Last &
DE Seeligeriolysin precursor (7
GN Name=180;
OS Listeria seeligeri.
OC Bacteria; Firmicutes; Bacill
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                                                                                                                                                                                                                                                                                                                                                                                                "Comparative human/murine sequence region 1 from human 3p21.3."; Mamm. Genome 13:646-655(2002).
EMBL; AJ428064; CAD20986.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAR-2002 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
01-OCT-2003 (TrEMBLrel.
Transmembrane protein 7.
                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                     MGD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
STRAIN=BALB/c; TISSUE=Liver;
MEDLINE=22350095; PubMed=12461651;
Kiss H., Darai E., Kiss C., Kost-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=Tmem7
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GO:0016021; C:integral to membrane;
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                                                                                                                                                                                                                                RVTVSVSQPSRNRKNYKVQV-----KIQNPT-----ACTANGSCDPSVTRQ----
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rsor (Thiol-activated cytolysin)
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 Bacillales; Listeriaceae;
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Pred. No. 0.
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Pred. No. 0.0028;
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Sciurognathi; Muridae;
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  Listeria.
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                                                                                                                                                            ALLSSPLYV 361
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SITE
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR001869; Thiol_cytolysin.

Pfam; PF01289; Thiol_cytolysin; 1.

PRINTS; PR01400; TACYTOLYSIN.

ProDom; PD007062; Thiol_cytolysin; 1.

PROSITE; PS00481; THIOL_CYTOLYSINS; 1.

Cytolysis; Hemolysis; Lipid-binding; Signal; Toxin.

SIGNAL 1

25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-92182018; PubMed=1543752;
Hass A., Dumbsky M., Kreft J.;
"Listeriolysin genes: complete sequence of ilo from Listeria ivanovii and of lso from Listeria seeligeri.";
Biochim. Biophys. Acta 1130:81-84(1992).
-!- FUNCTION: Sulfhydry1-activated toxin. Is able to lyse cholesterol containing membranes. Can be reversibly inactivated by oxidation. Cholesterol is the receptor for the binding of these toxins to eukaryotic cell membranes.
-!- SIMILARITY: Belongs to the thiol-activated cytolysin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; X60462; CAA42996.1; -. PIR; S22340; S22340. HSSP; P19995; 1PFO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=1640,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                209
                                                                                                                                                        154 TLSVDLÞGMTKKDNKIFVK--NÞTKSNVNNAVNTLVERWNDKYSKAYFNINAKIDYS--- 208
                                                                                                               99 YSTDEERAFVRTELAA 114
                                                                                                                                                                             49 TV$V$QP$ENRKNYKVQVKIQNPTACTANG$CDP$VTR--QKY------ADVTF$FTQ 98
                                                                                                                                                                                                                                         94 KDGSEYIVVEKKKKGINQNNADISVINAISSLTYPGALVKANRELVENQPNVLPVKRDSL 153
                                                                                                                                                                                                                                                                              13 KDGRQTLVL--NPRGVNPT-----NGVASLSQAGA------VPALEKRV 48
                                                                                                                                                                                                                                                                                                                       34;
                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                              ----DEMAYSESQLIA 220
                                                                                                                                                                                                                                                                                                                                                                                             1 25
26 530 Seeligeriolysin.
485 485 Binding to cholesterol (By similarity).
530 AA; 59181 MW; 416F7A4DD2029866 CRC64;
                                                                                                                                                                                                                                                                                                                     13.5%; Score 89.5; DB 1; Length 530; llarity 25.0%; Pred. No. 5.2; Conservative 22; Mismatches 37; Indels 4
                                                                                                                                                                                                                                                                                                                     43; Gaps
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